

PTO/SB/05 (2/98)

**UTILITY
PATENT APPLICATION
TRANSMITTAL**

(Only for nonprovisional applications under 37 CFR § 1.53(b))

Attorney Docket No.

210121.478C10

First Inventor or Application Identifier

Tongtong Wang

Title

COMPOSITIONS AND METHODS FOR THE THERAPY AND DIAGNOSIS OF LUNG CANCER

Express Mail Label No.

EL6125232081 US

APPLICATION ELEMENTS

See MPEP chapter 600 concerning utility patent application contents.

ADDRESS TO:

Box Patent Application
Assistant Commissioner for Patents
Washington, D.C. 20231

- | | |
|--|--|
| <p>1. <input type="checkbox"/> General Authorization Form & Fee Transmittal (Submit an original and a duplicate for fee processing)</p> <p>2. <input checked="" type="checkbox"/> Specification [Total Pages] 187 (preferred arrangement set forth below)</p> <ul style="list-style-type: none"> - Descriptive Title of the Invention - Cross References to Related Applications - Statement Regarding Fed sponsored R & D - Reference to Microfiche Appendix - Background of the Invention - Brief Summary of the Invention - Brief Description of the Drawings (if filed) - Detailed Description - Claim(s) - Abstract of the Disclosure <p>3. <input type="checkbox"/> Drawing(s) (35 USC 113) [Total Sheets] </p> <p>4. Oath or Declaration [Total Pages] </p> <p>a. <input type="checkbox"/> Newly executed (original or copy)</p> <p>b. <input type="checkbox"/> Copy from a prior application (37 CFR 1.63(d)) (for continuation/divisional with Box 17 completed)</p> <p style="margin-left: 20px;">i. <input type="checkbox"/> <u>DELETION OF INVENTOR(S)</u> Signed statement attached deleting inventor(s) named in the prior application, see 37 CFR 1.63(d)(2) and 1.33(b)</p> <p>5. <input type="checkbox"/> Incorporation By Reference (useable if box 4b is checked) The entire disclosure of the prior application, from which a copy of the oath or declaration is supplied under Box 4b, is considered to be part of the disclosure of the accompanying application and is hereby incorporated by reference therein.</p> | <p>6. <input type="checkbox"/> Microfiche Computer Program (Appendix)</p> <p>7. Nucleotide and Amino Acid Sequence Submission (if applicable, all necessary)</p> <p>a. <input checked="" type="checkbox"/> Computer-Readable Copy</p> <p>b. <input checked="" type="checkbox"/> Paper Copy (identical to computer copy)</p> <p>c. <input checked="" type="checkbox"/> Statement verifying identity of above copies</p> |
|--|--|

ACCOMPANYING APPLICATION PARTS

| | |
|---|--|
| <p>8. <input type="checkbox"/> Assignment Papers (cover sheet & document(s))</p> <p>9. <input type="checkbox"/> 37 CFR 3.73(b) Statement (when there is an assignee)</p> <p>10. <input type="checkbox"/> English Translation Document (if applicable)</p> <p>11. <input type="checkbox"/> Information Disclosure Statement (IDS)/PTO-1449</p> <p>12. <input type="checkbox"/> Preliminary Amendment</p> <p>13. <input checked="" type="checkbox"/> Return Receipt Postcard</p> <p>14. <input type="checkbox"/> Small Entity Statement(s)</p> <p>15. <input type="checkbox"/> Certified Copy of Priority Document(s) (if foreign priority is claimed)</p> <p>16. <input checked="" type="checkbox"/> Other: <u>Certificate of Express Mail</u></p> | <p><input type="checkbox"/> Power of Attorney</p> <p><input type="checkbox"/> Copies of IDS Citations</p> <p><input type="checkbox"/> Statement filed in prior application Status still proper and desired</p> |
|---|--|

ACCOMPANYING APPLICATION PARTS

8. ☐ Assignment Papers (cover sheet & document(s))
9. ☐ 37 CFR 3.73(b) Statement ☐ Power of Attorney
(when there is an assignee)
10. ☐ English Translation Document (if applicable)
11. ☐ Information Disclosure ☐ Copies of IDS
Statement (IDS)/PTO-1449 Citations
12. ☐ Preliminary Amendment
13. ☒ Return Receipt Postcard
14. ☐ Small Entity ☐ Statement filed in prior application,
Statement(s) Status still proper and desired
15. ☐ Certified Copy of Priority Document(s)
(if foreign priority is claimed)
16. ☒ Other: Certificate of Express Mail

17. If a CONTINUING APPLICATION, check appropriate box and supply the requisite information below and in a preliminary amendment

☐ Continuation ☐ Divisional ☒ Continuation-In-Part (CIP) of prior Application No.: **09/614,124**

Prior application information: Examiner not assigned

Group / Art Unit **not assigned**

☐ Claims the benefit of Provisional Application No.

CORRESPONDENCE ADDRESS

Jane E. R. Potter
Seed Intellectual Property Law Group PLLC
701 Fifth Avenue, Suite 6300
Seattle, Washington 98104-7092
Phone: (206) 622-4900 Fax: (206) 682-6031

Respectfully submitted,

TYPED or PRINTED NAME Jane E. R. Potter

SIGNATURE

u:\sharons\210121\478

REGISTRATION NO. 33.332

Date _____

August 29, 2008

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

PATENT

Applicants : Tongtong Wang, Medina, WA; Chaitanya S. Bangur, Seattle, WA;
Michael J. Lodes, Seattle, WA; Gary R. Fanger, Mill Creek, WA;
Thomas S. Vedvick, Federal Way, WA; Darrick Carter, Seattle, WA;
Marc W. Retter, Carnation, WA; Jane Mannion, Edmonds, WA;
Liquan Fan, Bellevue, WA

Filed : August 29, 2000

For : COMPOSITIONS AND METHODS FOR THE THERAPY AND
DIAGNOSIS OF LUNG CANCER



Docket No. : 210121.478C10
Date : August 29, 2000

Box Patent Application
Assistant Commissioner for Patents
Washington, DC 20231

CERTIFICATE OF MAILING BY "EXPRESS MAIL"

Assistant Commissioner for Patents:

I hereby certify that the enclosures listed below are being deposited with the United States Postal Service "EXPRESS MAIL Post Office to Addressee" service under 37 C.F.R. § 1.10, Mailing Label Certificate No. EL615232081US, on August 29, 2000, addressed to Box Patent Application, Assistant Commissioner for Patents, Washington, DC 20231.

Respectfully submitted,
Seed Intellectual Property Law Group PLLC

Steve Plante/Jeanette West/Susan Johnson

JEP:sds

Enclosures:

- Postcard
- Form PTO/SB/05
- Specification, Claims, Abstract (187 pages)
- Sequence Listing (518 pages)
- Declaration for Sequence Listing
- Diskette for Sequence Listing

u:\sharons\210121\478

COMPOSITIONS AND METHODS FOR THE THERAPY AND DIAGNOSIS OF LUNG CANCER

REFERENCE TO RELATED APPLICATIONS

This application is a continuation-in-part of U.S. Patent Application
5 No. 09/614,124, filed July 11, 2000, which is a continuation-in-part of U.S. Patent
Application No. 09/589,184, filed June 5, 2000, which is a continuation-in-part of U.S.
Patent Application No. 09/560,406, filed April 27, 2000, which is a continuation-in-part of
U.S. Patent Application No. 09/546,259, filed April 10, 2000, which is a continuation-in-
part of U.S. Patent Application No. 09/533,077, filed March 22, 2000, which is a
10 continuation-in-part of U.S. Patent Application No. 09/519,642 filed March 6, 2000, which
is a continuation-in-part of U.S. Patent Application No. 09/476,300, filed December 30,
1999, which is a continuation-in-part of U.S. Patent Application No. 09/466,867, filed
December 17, 1999, which is a continuation-in-part of U.S. Patent Application 09/419,356,
filed October 15, 1999, which is a continuation-in-part of U.S. Patent Application
15 No. 09/346,492, filed June 30, 1999, and is related to PCT/US00/18061, filed 6/30/00.

TECHNICAL FIELD OF THE INVENTION

The present invention relates generally to therapy and diagnosis of cancer,
such as lung cancer. The invention is more specifically related to polypeptides comprising
at least a portion of a lung tumor protein, and to polynucleotides encoding such
20 polypeptides. Such polypeptides and polynucleotides may be used in compositions for
prevention and treatment of lung cancer, and for the diagnosis and monitoring of such
cancers.

BACKGROUND OF THE INVENTION

Cancer is a significant health problem throughout the world. Although
25 advances have been made in detection and therapy of cancer, no vaccine or other
universally successful method for prevention or treatment is currently available. Current

therapies, which are generally based on a combination of chemotherapy or surgery and radiation, continue to prove inadequate in many patients.

Lung cancer is the primary cause of cancer death among both men and women in the U.S., with an estimated 172,000 new cases being reported in 1994. The five-
5 year survival rate among all lung cancer patients, regardless of the stage of disease at diagnosis, is only 13%. This contrasts with a five-year survival rate of 46% among cases detected while the disease is still localized. However, only 16% of lung cancers are discovered before the disease has spread.

Early detection is difficult since clinical symptoms are often not seen until
10 the disease has reached an advanced stage. Currently, diagnosis is aided by the use of chest x-rays, analysis of the type of cells contained in sputum and fiberoptic examination of the bronchial passages. Treatment regimens are determined by the type and stage of the cancer, and include surgery, radiation therapy and/or chemotherapy.

In spite of considerable research into therapies for this and other cancers,
15 lung cancer remains difficult to diagnose and treat effectively. Accordingly, there is a need in the art for improved methods for detecting and treating such cancers. The present invention fulfills these needs and further provides other related advantages.

SUMMARY OF THE INVENTION

Briefly stated, the present invention provides compositions and methods for
20 the diagnosis and therapy of cancer, such as lung cancer. In one aspect, the present invention provides polypeptides comprising at least a portion of a lung tumor protein, or a variant thereof. Certain portions and other variants are immunogenic, such that the ability of the variant to react with antigen-specific antisera is not substantially diminished. Within certain embodiments, the polypeptide comprises a sequence that is encoded by a
25 polynucleotide sequence selected from the group consisting of: (a) sequences recited in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180,

181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808 and 810-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 5 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, and 1669; (b) variants of a sequence recited in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 10 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 15 808, 810-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, and 1669; and 20 (c) complements of a sequence of (a) or (b). In specific embodiments, the polypeptides of the present invention comprise at least a portion of a tumor protein that includes an amino acid sequence selected from the group consisting of sequences recited in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809, 827, 1670-1675 and 1677-1678 and variants thereof.

25 The present invention further provides polynucleotides that encode a polypeptide as described above, or a portion thereof (such as a portion encoding at least 15 amino acid residues of a lung tumor protein), expression vectors comprising such polynucleotides and host cells transformed or transfected with such expression vectors.

Within other aspects, the present invention provides pharmaceutical compositions comprising a polypeptide or polynucleotide as described above and a physiologically acceptable carrier.

Within a related aspect of the present invention, vaccines, or immunogenic
5 compositions, for prophylactic or therapeutic use are provided. Such vaccines comprise a polypeptide or polynucleotide as described above and an immunostimulant.

The present invention further provides pharmaceutical compositions that comprise: (a) an antibody or antigen-binding fragment thereof that specifically binds to a lung tumor protein; and (b) a physiologically acceptable carrier.

10 Within further aspects, the present invention provides pharmaceutical compositions comprising: (a) an antigen presenting cell that expresses a polypeptide as described above and (b) a pharmaceutically acceptable carrier or excipient. Antigen presenting cells include dendritic cells, macrophages, monocytes, fibroblasts and B cells.

Within related aspects, vaccines, or immunogenic compositions, are
15 provided that comprise: (a) an antigen presenting cell that expresses a polypeptide as described above and (b) an immunostimulant.

The present invention further provides, in other aspects, fusion proteins that comprise at least one polypeptide as described above, as well as polynucleotides encoding such fusion proteins.

20 Within related aspects, pharmaceutical compositions comprising a fusion protein, or a polynucleotide encoding a fusion protein, in combination with a physiologically acceptable carrier are provided.

Vaccines, or immunogenic compositions, are further provided, within other aspects, that comprise a fusion protein, or a polynucleotide encoding a fusion protein, in
25 combination with an immunostimulant.

Within further aspects, the present invention provides methods for inhibiting the development of a cancer in a patient, comprising administering to a patient a pharmaceutical composition or immunogenic composition as recited above. The patient

may be afflicted with lung cancer, in which case the methods provide treatment for the disease, or patient considered at risk for such a disease may be treated prophylactically.

The present invention further provides, within other aspects, methods for removing tumor cells from a biological sample, comprising contacting a biological sample
5 with T cells that specifically react with a lung tumor protein, wherein the step of contacting is performed under conditions and for a time sufficient to permit the removal of cells expressing the protein from the sample.

Within related aspects, methods are provided for inhibiting the development of a cancer in a patient, comprising administering to a patient a biological sample treated as
10 described above.

Methods are further provided, within other aspects, for stimulating and/or expanding T cells specific for a lung tumor protein, comprising contacting T cells with one or more of: (i) a polypeptide as described above; (ii) a polynucleotide encoding such a polypeptide; and/or (iii) an antigen presenting cell that expresses such a polypeptide; under
15 conditions and for a time sufficient to permit the stimulation and/or expansion of T cells. Isolated T cell populations comprising T cells prepared as described above are also provided.

Within further aspects, the present invention provides methods for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective
20 amount of a T cell population as described above.

The present invention further provides methods for inhibiting the development of a cancer in a patient, comprising the steps of: (a) incubating CD4⁺ and/or CD8⁺ T cells isolated from a patient with one or more of: (i) a polypeptide comprising at least an immunogenic portion of a lung tumor protein; (ii) a polynucleotide encoding such
25 a polypeptide; and (iii) an antigen-presenting cell that expressed such a polypeptide; and (b) administering to the patient an effective amount of the proliferated T cells, and thereby inhibiting the development of a cancer in the patient. Proliferated cells may, but need not, be cloned prior to administration to the patient.

Within further aspects, the present invention provides methods for determining the presence or absence of a cancer in a patient, comprising: (a) contacting a biological sample obtained from a patient with a binding agent that binds to a polypeptide as recited above; (b) detecting in the sample an amount of polypeptide that binds to the binding agent; and (c) comparing the amount of polypeptide with a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient. Within preferred embodiments, the binding agent is an antibody, more preferably a monoclonal antibody. The cancer may be lung cancer.

The present invention also provides, within other aspects, methods for monitoring the progression of a cancer in a patient. Such methods comprise the steps of: (a) contacting a biological sample obtained from a patient at a first point in time with a binding agent that binds to a polypeptide as recited above; (b) detecting in the sample an amount of polypeptide that binds to the binding agent; (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and (d) comparing the amount of polypeptide detected in step (c) with the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

The present invention further provides, within other aspects, methods for determining the presence or absence of a cancer in a patient, comprising the steps of: (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein; (b) detecting in the sample a level of a polynucleotide, preferably mRNA, that hybridizes to the oligonucleotide; and (c) comparing the level of polynucleotide that hybridizes to the oligonucleotide with a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient. Within certain embodiments, the amount of mRNA is detected via polymerase chain reaction using, for example, at least one oligonucleotide primer that hybridizes to a polynucleotide encoding a polypeptide as recited above, or a complement of such a polynucleotide. Within other embodiments, the amount of mRNA is detected using a hybridization technique, employing an

oligonucleotide probe that hybridizes to a polynucleotide that encodes a polypeptide as recited above, or a complement of such a polynucleotide.

In related aspects, methods are provided for monitoring the progression of a cancer in a patient, comprising the steps of: (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein; (b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide; (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and (d) comparing the amount of polynucleotide detected in step (c) with the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

Within further aspects, the present invention provides antibodies, such as monoclonal antibodies, that bind to a polypeptide as described above, as well as diagnostic kits comprising such antibodies. Diagnostic kits comprising one or more oligonucleotide probes or primers as described above are also provided.

These and other aspects of the present invention will become apparent upon reference to the following detailed description. All references disclosed herein are hereby incorporated by reference in their entirety as if each was incorporated individually.

SEQUENCE IDENTIFIERS

SEQ ID NO: 1 is the determined cDNA sequence for clone #19038, also referred to as L845P.

SEQ ID NO: 2 is the determined cDNA sequence for clone #19036.

SEQ ID NO: 3 is the determined cDNA sequence for clone #19034.

SEQ ID NO: 4 is the determined cDNA sequence for clone #19033.

SEQ ID NO: 5 is the determined cDNA sequence for clone #19032.

SEQ ID NO: 6 is the determined cDNA sequence for clone #19030, also referred to as L559S.

SEQ ID NO: 7 is the determined cDNA sequence for clone #19029.

SEQ ID NO: 8 is the determined cDNA sequence for clone #19025.

5

19007, 19016 and 19017.

10

15

SEQ ID NO: 14 is the determined cDNA sequence for clone #19004.

10

15

SEQ ID NO: 21 is the determined cDNA sequence for clone #18992.

20

SEQ ID NO: 25 is the determined cDNA sequence for clone #18985, also referred as L839P.

25

SEQ ID NO: 26 is the determined cDNA sequence for clone #18984, also referred to as L847P.

25

SEQ ID NO: 28 is the determined cDNA sequence for clones #18976 and 18980.

SEQ ID NO: 29 is the determined cDNA sequence for clone #18975.

SEQ ID NO: 30 is the determined cDNA sequence for clone #18974.

SEQ ID NO: 31 is the determined cDNA sequence for clone #18973.

SEQ ID NO: 32 is the determined cDNA sequence for clone #18972.

SEQ ID NO: 33 is the determined cDNA sequence for clone #18971, also referred to as L801P.

5 SEQ ID NO: 34 is the determined cDNA sequence for clone #18970.

SEQ ID NO: 35 is the determined cDNA sequence for clone #18966.

SEQ ID NO: 36 is the determined cDNA sequence for clones #18964, 18968 and 19039.

SEQ ID NO: 37 is the determined cDNA sequence for clone #18960.

10 SEQ ID NO: 38 is the determined cDNA sequence for clone #18959.

SEQ ID NO: 39 is the determined cDNA sequence for clones #18958 and 18982.

SEQ ID NO: 40 is the determined cDNA sequence for clones #18956 and 19015.

15 SEQ ID NO: 41 is the determined cDNA sequence for clone #18954, also referred to L848P.

SEQ ID NO: 42 is the determined cDNA sequence for clone #18951.

SEQ ID NO: 43 is the determined cDNA sequence for clone #18950.

20 SEQ ID NO: 44 is the determined cDNA sequence for clones #18949 and 19024, also referred to as L844P.

SEQ ID NO: 45 is the determined cDNA sequence for clone #18948.

SEQ ID NO: 46 is the determined cDNA sequence for clone #18947, also referred to as L840P.

25 SEQ ID NO: 47 is the determined cDNA sequence for clones #18946, 18953, 18969 and 19027.

SEQ ID NO: 48 is the determined cDNA sequence for clone #18942.

SEQ ID NO: 49 is the determined cDNA sequence for clone #18940, 18962, 18963, 19006, 19008, 19000, and 19031.

SEQ ID NO: 50 is the determined cDNA sequence for clone #18939.

SEQ ID NO: 51 is the determined cDNA sequence for clones #18938 and 18952.

SEQ ID NO: 52 is the determined cDNA sequence for clone #18938.

SEQ ID NO: 53 is the determined cDNA sequence for clone #18937.

5 SEQ ID NO: 54 is the determined cDNA sequence for clones #18934, 18935, 18993 and 19022, also referred to as L548S.

SEQ ID NO: 55 is the determined cDNA sequence for clone #18932.

SEQ ID NO: 56 is the determined cDNA sequence for clones #18931 and 18936.

10 SEQ ID NO: 57 is the determined cDNA sequence for clone #18930.

SEQ ID NO: 58 is the determined cDNA sequence for clone #19014, also referred to as L773P.

SEQ ID NO: 59 is the determined cDNA sequence for clone #19127.

15 SEQ ID NO: 60 is the determined cDNA sequence for clones #19057 and 19064.

SEQ ID NO: 61 is the determined cDNA sequence for clone #19122.

SEQ ID NO: 62 is the determined cDNA sequence for clones #19120 and 18121.

SEQ ID NO: 63 is the determined cDNA sequence for clone #19118.

20 SEQ ID NO: 64 is the determined cDNA sequence for clone #19117.

SEQ ID NO: 65 is the determined cDNA sequence for clone #19116.

SEQ ID NO: 66 is the determined cDNA sequence for clone #19114.

SEQ ID NO: 67 is the determined cDNA sequence for clone #19112, also known as L561S.

25 SEQ ID NO: 68 is the determined cDNA sequence for clone #19110.

SEQ ID NO: 69 is the determined cDNA sequence for clone #19107, also referred to as L552S.

SEQ ID NO: 70 is the determined cDNA sequence for clone #19106, also referred to as L547S.

SEQ ID NO: 71 is the determined cDNA sequence for clones #19105 and 19111.

SEQ ID NO: 72 is the determined cDNA sequence for clone #19099.

SEQ ID NO: 73 is the determined cDNA sequence for clones #19095,
5 19104 and 19125, also referred to as L549S.

SEQ ID NO: 74 is the determined cDNA sequence for clone #19094.

SEQ ID NO: 75 is the determined cDNA sequence for clones #19089 and 19101.

SEQ ID NO: 76 is the determined cDNA sequence for clone #19088.

10 SEQ ID NO: 77 is the determined cDNA sequence for clones #19087, 19092, 19096, 19100 and 19119.

SEQ ID NO: 78 is the determined cDNA sequence for clone #19086.

SEQ ID NO: 79 is the determined cDNA sequence for clone #19085, also referred to as L550S.

15 SEQ ID NO: 80 is the determined cDNA sequence for clone #19084, also referred to as clone #19079.

SEQ ID NO: 81 is the determined cDNA sequence for clone #19082.

SEQ ID NO: 82 is the determined cDNA sequence for clone #19080.

SEQ ID NO: 83 is the determined cDNA sequence for clone #19077.

20 SEQ ID NO: 84 is the determined cDNA sequence for clone #19076, also referred to as L551S.

SEQ ID NO: 85 is the determined cDNA sequence for clone #19074, also referred to as clone #20102.

25 SEQ ID NO: 86 is the determined cDNA sequence for clone #19073, also referred to as L560S.

SEQ ID NO: 87 is the determined cDNA sequence for clones #19072 and 19115.

SEQ ID NO: 88 is the determined cDNA sequence for clone #19071.

SEQ ID NO: 89 is the determined cDNA sequence for clone #19070.

SEQ ID NO: 90 is the determined cDNA sequence for clone #19069.

SEQ ID NO: 91 is the determined cDNA sequence for clone #19068, also referred to L563S.

SEQ ID NO: 92 is the determined cDNA sequence for clone #19066.

5 SEQ ID NO: 93 is the determined cDNA sequence for clone #19065.

SEQ ID NO: 94 is the determined cDNA sequence for clone #19063.

SEQ ID NO: 95 is the determined cDNA sequence for clones #19061, 19081, 19108 and 19109.

10 SEQ ID NO: 96 is the determined cDNA sequence for clones #19060, 19067 and 19083, also referred to as L548S.

SEQ ID NO: 97 is the determined cDNA sequence for clones #19059 and 19062.

SEQ ID NO: 98 is the determined cDNA sequence for clone #19058.

SEQ ID NO: 99 is the determined cDNA sequence for clone #19124.

15 SEQ ID NO: 100 is the determined cDNA sequence for clone #18929.

SEQ ID NO: 101 is the determined cDNA sequence for clone #18422.

SEQ ID NO: 102 is the determined cDNA sequence for clone #18425.

SEQ ID NO: 103 is the determined cDNA sequence for clone #18431.

SEQ ID NO: 104 is the determined cDNA sequence for clone #18433.

20 SEQ ID NO: 105 is the determined cDNA sequence for clone #18444.

SEQ ID NO: 106 is the determined cDNA sequence for clone #18449.

SEQ ID NO: 107 is the determined cDNA sequence for clone #18451.

SEQ ID NO: 108 is the determined cDNA sequence for clone #18452.

SEQ ID NO: 109 is the determined cDNA sequence for clone #18455.

25 SEQ ID NO: 110 is the determined cDNA sequence for clone #18457.

SEQ ID NO: 111 is the determined cDNA sequence for clone #18466.

SEQ ID NO: 112 is the determined cDNA sequence for clone #18468.

SEQ ID NO: 113 is the determined cDNA sequence for clone #18471.

SEQ ID NO: 114 is the determined cDNA sequence for clone #18475.

SEQ ID NO: 115 is the determined cDNA sequence for clone #18476.
 SEQ ID NO: 116 is the determined cDNA sequence for clone #18477.
 SEQ ID NO: 117 is the determined cDNA sequence for clone #20631.
 SEQ ID NO: 118 is the determined cDNA sequence for clone #20634.
 5 SEQ ID NO: 119 is the determined cDNA sequence for clone #20635.
 SEQ ID NO: 120 is the determined cDNA sequence for clone #20637.
 SEQ ID NO: 121 is the determined cDNA sequence for clone #20638.
 SEQ ID NO: 122 is the determined cDNA sequence for clone #20643.
 SEQ ID NO: 123 is the determined cDNA sequence for clone #20652.
 10 SEQ ID NO: 124 is the determined cDNA sequence for clone #20653.
 SEQ ID NO: 125 is the determined cDNA sequence for clone #20657.
 SEQ ID NO: 126 is the determined cDNA sequence for clone #20658.
 SEQ ID NO: 127 is the determined cDNA sequence for clone #20660.
 SEQ ID NO: 128 is the determined cDNA sequence for clone #20661.
 15 SEQ ID NO: 129 is the determined cDNA sequence for clone #20663.
 SEQ ID NO: 130 is the determined cDNA sequence for clone #20665.
 SEQ ID NO: 131 is the determined cDNA sequence for clone #20670.
 SEQ ID NO: 132 is the determined cDNA sequence for clone #20671.
 SEQ ID NO: 133 is the determined cDNA sequence for clone #20672.
 20 SEQ ID NO: 134 is the determined cDNA sequence for clone #20675.
 SEQ ID NO: 135 is the determined cDNA sequence for clone #20679.
 SEQ ID NO: 136 is the determined cDNA sequence for clone #20681.
 SEQ ID NO: 137 is the determined cDNA sequence for clone #20682.
 SEQ ID NO: 138 is the determined cDNA sequence for clone #20684.
 25 SEQ ID NO: 139 is the determined cDNA sequence for clone #20685.
 SEQ ID NO: 140 is the determined cDNA sequence for clone #20689.
 SEQ ID NO: 141 is the determined cDNA sequence for clone #20699.
 SEQ ID NO: 142 is the determined cDNA sequence for clone #20701.
 SEQ ID NO: 143 is the determined cDNA sequence for clone #20702.

SEQ ID NO: 144 is the determined cDNA sequence for clone #20708.
 SEQ ID NO: 145 is the determined cDNA sequence for clone #20715.
 SEQ ID NO: 146 is the determined cDNA sequence for clone #20716.
 SEQ ID NO: 147 is the determined cDNA sequence for clone #20719.
 5 SEQ ID NO: 148 is the determined cDNA sequence for clone #19129.
 SEQ ID NO: 149 is the determined cDNA sequence for clone #19131.1.
 SEQ ID NO: 150 is the determined cDNA sequence for clone #19132.2.
 SEQ ID NO: 151 is the determined cDNA sequence for clone #19133.
 SEQ ID NO: 152 is the determined cDNA sequence for clone #19134.2.
 10 SEQ ID NO: 153 is the determined cDNA sequence for clone #19135.2.
 SEQ ID NO: 154 is the determined cDNA sequence for clone #19137.
 SEQ ID NO: 155 is a first determined cDNA sequence for clone #19138.1.
 SEQ ID NO: 156 is a second determined cDNA sequence for clone
 #19138.2.
 15 SEQ ID NO: 157 is the determined cDNA sequence for clone #19139.
 SEQ ID NO: 158 is a first determined cDNA sequence for clone #19140.1.
 SEQ ID NO: 159 is a second determined cDNA sequence for clone
 #19140.2.
 20 SEQ ID NO: 160 is the determined cDNA sequence for clone #19141.
 SEQ ID NO: 161 is the determined cDNA sequence for clone #19143.
 SEQ ID NO: 162 is the determined cDNA sequence for clone #19144.
 SEQ ID NO: 163 is a first determined cDNA sequence for clone #19145.1.
 SEQ ID NO: 164 is a second determined cDNA sequence for clone
 #19145.2.
 25 SEQ ID NO: 165 is the determined cDNA sequence for clone #19146.
 SEQ ID NO: 166 is the determined cDNA sequence for clone #19149.1.
 SEQ ID NO: 167 is the determined cDNA sequence for clone #19152.
 SEQ ID NO: 168 is a first determined cDNA sequence for clone #19153.1.

SEQ ID NO: 169 is a second determined cDNA sequence for clone #19153.2.

SEQ ID NO: 170 is the determined cDNA sequence for clone #19155.

SEQ ID NO: 171 is the determined cDNA sequence for clone #19157.

5 SEQ ID NO: 172 is the determined cDNA sequence for clone #19159.

SEQ ID NO: 173 is the determined cDNA sequence for clone #19160.

SEQ ID NO: 174 is a first determined cDNA sequence for clone #19161.1.

SEQ ID NO: 175 is a second determined cDNA sequence for clone #19161.2.

10 SEQ ID NO: 176 is the determined cDNA sequence for clone #19162.1.

SEQ ID NO: 177 is the determined cDNA sequence for clone #19166.

SEQ ID NO: 178 is the determined cDNA sequence for clone #19169.

SEQ ID NO: 179 is the determined cDNA sequence for clone #19171.

SEQ ID NO: 180 is a first determined cDNA sequence for clone #19173.1.

15 SEQ ID NO: 181 is a second determined cDNA sequence for clone #19173.2.

SEQ ID NO: 182 is the determined cDNA sequence for clone #19174.1.

SEQ ID NO: 183 is the determined cDNA sequence for clone #19175.

SEQ ID NO: 184 is the determined cDNA sequence for clone #19177.

20 SEQ ID NO: 185 is the determined cDNA sequence for clone #19178.

SEQ ID NO: 186 is the determined cDNA sequence for clone #19179.1.

SEQ ID NO: 187 is the determined cDNA sequence for clone #19179.2.

SEQ ID NO: 188 is the determined cDNA sequence for clone #19180.

SEQ ID NO: 189 is a first determined cDNA sequence for clone #19182.1.

25 SEQ ID NO: 190 is a second determined cDNA sequence for clone #19182.2.

SEQ ID NO: 191 is the determined cDNA sequence for clone #19183.1.

SEQ ID NO: 192 is the determined cDNA sequence for clone #19185.1.

SEQ ID NO: 193 is the determined cDNA sequence for clone #19187.

SEQ ID NO: 194 is the determined cDNA sequence for clone #19188.
 SEQ ID NO: 195 is the determined cDNA sequence for clone #19190.
 SEQ ID NO: 196 is the determined cDNA sequence for clone #19191.
 SEQ ID NO: 197 is the determined cDNA sequence for clone #19192.
 5 SEQ ID NO: 198 is the determined cDNA sequence for clone #19193.
 SEQ ID NO: 199 is a first determined cDNA sequence for clone #19194.1.
 SEQ ID NO: 200 is a second determined cDNA sequence for clone
 #19194.2.
 SEQ ID NO: 201 is the determined cDNA sequence for clone #19197.
 10 SEQ ID NO: 202 is a first determined cDNA sequence for clone #19200.1.
 SEQ ID NO: 203 is a second determined cDNA sequence for clone
 #19200.2.
 SEQ ID NO: 204 is the determined cDNA sequence for clone #19202.
 SEQ ID NO: 205 is a first determined cDNA sequence for clone #19204.1.
 15 SEQ ID NO: 206 is a second determined cDNA sequence for clone
 #19204.2.
 SEQ ID NO: 207 is the determined cDNA sequence for clone #19205.
 SEQ ID NO: 208 is a first determined cDNA sequence for clone #19206.1.
 SEQ ID NO: 209 is a second determined cDNA sequence for clone
 20 #19206.2.
 SEQ ID NO: 210 is the determined cDNA sequence for clone #19207.
 SEQ ID NO: 211 is the determined cDNA sequence for clone #19208.
 SEQ ID NO: 212 is a first determined cDNA sequence for clone #19211.1.
 SEQ ID NO: 213 is a second determined cDNA sequence for clone
 25 #19211.2.
 SEQ ID NO: 214 is a first determined cDNA sequence for clone #19214.1.
 SEQ ID NO: 215 is a second determined cDNA sequence for clone
 #19214.2.
 SEQ ID NO: 216 is the determined cDNA sequence for clone #19215.

SEQ ID NO: 217 is a first determined cDNA sequence for clone #19217. 2.
 SEQ ID NO: 218 is a second determined cDNA sequence for clone
 #19217.2.

5 SEQ ID NO: 219 is a first determined cDNA sequence for clone #19218.1.
 SEQ ID NO: 220 is a second determined cDNA sequence for clone
 #19218.2.

SEQ ID NO: 221 is a first determined cDNA sequence for clone #19220.1.
 SEQ ID NO: 222 is a second determined cDNA sequence for clone
 #19220.2.

10 SEQ ID NO: 223 is the determined cDNA sequence for clone #22015.
 SEQ ID NO: 224 is the determined cDNA sequence for clone #22017.
 SEQ ID NO: 225 is the determined cDNA sequence for clone #22019.
 SEQ ID NO: 226 is the determined cDNA sequence for clone #22020.
 SEQ ID NO: 227 is the determined cDNA sequence for clone #22023.

15 SEQ ID NO: 228 is the determined cDNA sequence for clone #22026.
 SEQ ID NO: 229 is the determined cDNA sequence for clone #22027.
 SEQ ID NO: 230 is the determined cDNA sequence for clone #22028.
 SEQ ID NO: 231 is the determined cDNA sequence for clone #22032.
 SEQ ID NO: 232 is the determined cDNA sequence for clone #22037.

20 SEQ ID NO: 233 is the determined cDNA sequence for clone #22045.
 SEQ ID NO: 234 is the determined cDNA sequence for clone #22048.
 SEQ ID NO: 235 is the determined cDNA sequence for clone #22050.
 SEQ ID NO: 236 is the determined cDNA sequence for clone #22052.
 SEQ ID NO: 237 is the determined cDNA sequence for clone #22053.

25 SEQ ID NO: 238 is the determined cDNA sequence for clone #22057.
 SEQ ID NO: 239 is the determined cDNA sequence for clone #22066.
 SEQ ID NO: 240 is the determined cDNA sequence for clone #22077.
 SEQ ID NO: 241 is the determined cDNA sequence for clone #22085.
 SEQ ID NO: 242 is the determined cDNA sequence for clone #22105.

SEQ ID NO: 243 is the determined cDNA sequence for clone #22108.
 SEQ ID NO: 244 is the determined cDNA sequence for clone #22109.
 SEQ ID NO: 245 is the determined cDNA sequence for clone #24842.
 SEQ ID NO: 246 is the determined cDNA sequence for clone #24843.
 5 SEQ ID NO: 247 is the determined cDNA sequence for clone #24845.
 SEQ ID NO: 248 is the determined cDNA sequence for clone #24851.
 SEQ ID NO: 249 is the determined cDNA sequence for clone #24852.
 SEQ ID NO: 250 is the determined cDNA sequence for clone #24853.
 SEQ ID NO: 251 is the determined cDNA sequence for clone #24854.
 10 SEQ ID NO: 252 is the determined cDNA sequence for clone #24855.
 SEQ ID NO: 253 is the determined cDNA sequence for clone #24860.
 SEQ ID NO: 254 is the determined cDNA sequence for clone #24864.
 SEQ ID NO: 255 is the determined cDNA sequence for clone #24866.
 SEQ ID NO: 256 is the determined cDNA sequence for clone #24867.
 15 SEQ ID NO: 257 is the determined cDNA sequence for clone #24868.
 SEQ ID NO: 258 is the determined cDNA sequence for clone #24869.
 SEQ ID NO: 259 is the determined cDNA sequence for clone #24870.
 SEQ ID NO: 260 is the determined cDNA sequence for clone #24872.
 SEQ ID NO: 261 is the determined cDNA sequence for clone #24873.
 20 SEQ ID NO: 262 is the determined cDNA sequence for clone #24875.
 SEQ ID NO: 263 is the determined cDNA sequence for clone #24882.
 SEQ ID NO: 264 is the determined cDNA sequence for clone #24885.
 SEQ ID NO: 265 is the determined cDNA sequence for clone #24886.
 SEQ ID NO: 266 is the determined cDNA sequence for clone #24887.
 25 SEQ ID NO: 267 is the determined cDNA sequence for clone #24888.
 SEQ ID NO: 268 is the determined cDNA sequence for clone #24890.
 SEQ ID NO: 269 is the determined cDNA sequence for clone #24896.
 SEQ ID NO: 270 is the determined cDNA sequence for clone #24897.
 SEQ ID NO: 271 is the determined cDNA sequence for clone #24899.

SEQ ID NO: 272 is the determined cDNA sequence for clone #24901.
 SEQ ID NO: 273 is the determined cDNA sequence for clone #24902.
 SEQ ID NO: 274 is the determined cDNA sequence for clone #24906.
 SEQ ID NO: 275 is the determined cDNA sequence for clone #24912.
 5 SEQ ID NO: 276 is the determined cDNA sequence for clone #24913.
 SEQ ID NO: 277 is the determined cDNA sequence for clone #24920.
 SEQ ID NO: 278 is the determined cDNA sequence for clone #24927.
 SEQ ID NO: 279 is the determined cDNA sequence for clone #24930.
 SEQ ID NO: 280 is the determined cDNA sequence for clone #26938.
 10 SEQ ID NO: 281 is the determined cDNA sequence for clone #26939.
 SEQ ID NO: 282 is the determined cDNA sequence for clone #26943.
 SEQ ID NO: 283 is the determined cDNA sequence for clone #26948.
 SEQ ID NO: 284 is the determined cDNA sequence for clone #26951.
 SEQ ID NO: 285 is the determined cDNA sequence for clone #26955.
 15 SEQ ID NO: 286 is the determined cDNA sequence for clone #26956.
 SEQ ID NO: 287 is the determined cDNA sequence for clone #26959.
 SEQ ID NO: 288 is the determined cDNA sequence for clone #26961.
 SEQ ID NO: 289 is the determined cDNA sequence for clone #26962.
 SEQ ID NO: 290 is the determined cDNA sequence for clone #26964.
 20 SEQ ID NO: 291 is the determined cDNA sequence for clone #26966.
 SEQ ID NO: 292 is the determined cDNA sequence for clone #26968.
 SEQ ID NO: 293 is the determined cDNA sequence for clone #26972.
 SEQ ID NO: 294 is the determined cDNA sequence for clone #26973.
 SEQ ID NO: 295 is the determined cDNA sequence for clone #26974.
 25 SEQ ID NO: 296 is the determined cDNA sequence for clone #26976.
 SEQ ID NO: 297 is the determined cDNA sequence for clone #26977.
 SEQ ID NO: 298 is the determined cDNA sequence for clone #26979.
 SEQ ID NO: 299 is the determined cDNA sequence for clone #26980.
 SEQ ID NO: 300 is the determined cDNA sequence for clone #26981.

- SEQ ID NO: 301 is the determined cDNA sequence for clone #26984.
 SEQ ID NO: 302 is the determined cDNA sequence for clone #26985.
 SEQ ID NO: 303 is the determined cDNA sequence for clone #26986.
 SEQ ID NO: 304 is the determined cDNA sequence for clone #26993.
 5 SEQ ID NO: 305 is the determined cDNA sequence for clone #26994.
 SEQ ID NO: 306 is the determined cDNA sequence for clone #26995.
 SEQ ID NO: 307 is the determined cDNA sequence for clone #27003.
 SEQ ID NO: 308 is the determined cDNA sequence for clone #27005.
 SEQ ID NO: 309 is the determined cDNA sequence for clone #27010.
 10 SEQ ID NO: 310 is the determined cDNA sequence for clone #27011.
 SEQ ID NO: 311 is the determined cDNA sequence for clone #27013.
 SEQ ID NO: 312 is the determined cDNA sequence for clone #27016
 SEQ ID NO: 313 is the determined cDNA sequence for clone #27017.
 SEQ ID NO: 314 is the determined cDNA sequence for clone #27019.
 15 SEQ ID NO: 315 is the determined cDNA sequence for clone #27028.
 SEQ ID NO: 316 is the full-length cDNA sequence for clone #19060.
 SEQ ID NO: 317 is the full-length cDNA sequence for clone #18964.
 SEQ ID NO: 318 is the full-length cDNA sequence for clone #18929.
 SEQ ID NO: 319 is the full-length cDNA sequence for clone #18991.
 20 SEQ ID NO: 320 is the full-length cDNA sequence for clone #18996.
 SEQ ID NO: 321 is the full-length cDNA sequence for clone #18966.
 SEQ ID NO: 322 is the full-length cDNA sequence for clone #18951.
 SEQ ID NO: 323 is the full-length cDNA sequence for clone #18973 (also
 known as L516S).
 25 SEQ ID NO: 324 is the amino acid sequence for clone #19060.
 SEQ ID NO: 325 is the amino acid sequence for clone #19063.
 SEQ ID NO: 326 is the amino acid sequence for clone #19077.
 SEQ ID NO: 327 is the amino acid sequence for clone #19110.
 SEQ ID NO: 328 is the amino acid sequence for clone #19122.

5 SEQ ID NO: 329 is the amino acid sequence for clone #19118.
 SEQ ID NO: 330 is the amino acid sequence for clone #19080.
 SEQ ID NO: 331 is the amino acid sequence for clone #19127.
 SEQ ID NO: 332 is the amino acid sequence for clone #19117.
 SEQ ID NO: 333 is the amino acid sequence for clone #19095, also referred
 to L549S.

10 SEQ ID NO: 334 is the amino acid sequence for clone #18964.
 SEQ ID NO: 335 is the amino acid sequence for clone #18929.
 SEQ ID NO: 336 is the amino acid sequence for clone #18991.
 SEQ ID NO: 337 is the amino acid sequence for clone #18996.
 SEQ ID NO: 338 is the amino acid sequence for clone #18966.
 SEQ ID NO: 339 is the amino acid sequence for clone #18951.
 SEQ ID NO: 340 is the amino acid sequence for clone #18973.
 SEQ ID NO: 341 is the determined cDNA sequence for clone 26461.
 15 SEQ ID NO: 342 is the determined cDNA sequence for clone 26462.
 SEQ ID NO: 343 is the determined cDNA sequence for clone 26463.
 SEQ ID NO: 344 is the determined cDNA sequence for clone 26464.
 SEQ ID NO: 345 is the determined cDNA sequence for clone 26465.
 SEQ ID NO: 346 is the determined cDNA sequence for clone 26466.
 20 SEQ ID NO: 347 is the determined cDNA sequence for clone 26467.
 SEQ ID NO: 348 is the determined cDNA sequence for clone 26468.
 SEQ ID NO: 349 is the determined cDNA sequence for clone 26469.
 SEQ ID NO: 350 is the determined cDNA sequence for clone 26470.
 SEQ ID NO: 351 is the determined cDNA sequence for clone 26471.
 25 SEQ ID NO: 352 is the determined cDNA sequence for clone 26472.
 SEQ ID NO: 353 is the determined cDNA sequence for clone 26474.
 SEQ ID NO: 354 is the determined cDNA sequence for clone 26475.
 SEQ ID NO: 355 is the determined cDNA sequence for clone 26476.
 SEQ ID NO: 356 is the determined cDNA sequence for clone 26477.

SEQ ID NO: 357 is the determined cDNA sequence for clone 26478.
 SEQ ID NO: 358 is the determined cDNA sequence for clone 26479.
 SEQ ID NO: 359 is the determined cDNA sequence for clone 26480.
 SEQ ID NO: 360 is the determined cDNA sequence for clone 26481.
 5 SEQ ID NO: 361 is the determined cDNA sequence for clone 26482
 SEQ ID NO: 362 is the determined cDNA sequence for clone 26483.
 SEQ ID NO: 363 is the determined cDNA sequence for clone 26484.
 SEQ ID NO: 364 is the determined cDNA sequence for clone 26485.
 SEQ ID NO: 365 is the determined cDNA sequence for clone 26486.
 10 SEQ ID NO: 366 is the determined cDNA sequence for clone 26487.
 SEQ ID NO: 367 is the determined cDNA sequence for clone 26488.
 SEQ ID NO: 368 is the determined cDNA sequence for clone 26489.
 SEQ ID NO: 369 is the determined cDNA sequence for clone 26490.
 SEQ ID NO: 370 is the determined cDNA sequence for clone 26491.
 15 SEQ ID NO: 371 is the determined cDNA sequence for clone 26492.
 SEQ ID NO: 372 is the determined cDNA sequence for clone 26493.
 SEQ ID NO: 373 is the determined cDNA sequence for clone 26494.
 SEQ ID NO: 374 is the determined cDNA sequence for clone 26495.
 SEQ ID NO: 375 is the determined cDNA sequence for clone 26496.
 20 SEQ ID NO: 376 is the determined cDNA sequence for clone 26497.
 SEQ ID NO: 377 is the determined cDNA sequence for clone 26498.
 SEQ ID NO: 378 is the determined cDNA sequence for clone 26499.
 SEQ ID NO: 379 is the determined cDNA sequence for clone 26500.
 SEQ ID NO: 380 is the determined cDNA sequence for clone 26501.
 25 SEQ ID NO: 381 is the determined cDNA sequence for clone 26502.
 SEQ ID NO: 382 is the determined cDNA sequence for clone 26503.
 SEQ ID NO: 383 is the determined cDNA sequence for clone 26504.
 SEQ ID NO: 384 is the determined cDNA sequence for clone 26505.
 SEQ ID NO: 385 is the determined cDNA sequence for clone 26506.

SEQ ID NO: 386 is the determined cDNA sequence for clone 26507.
 SEQ ID NO: 387 is the determined cDNA sequence for clone 26508.
 SEQ ID NO: 388 is the determined cDNA sequence for clone 26509.
 SEQ ID NO: 389 is the determined cDNA sequence for clone 26511.
 5 SEQ ID NO: 390 is the determined cDNA sequence for clone 26513.
 SEQ ID NO: 391 is the determined cDNA sequence for clone 26514.
 SEQ ID NO: 392 is the determined cDNA sequence for clone 26515.
 SEQ ID NO: 393 is the determined cDNA sequence for clone 26516.
 SEQ ID NO: 394 is the determined cDNA sequence for clone 26517.
 10 SEQ ID NO: 395 is the determined cDNA sequence for clone 26518.
 SEQ ID NO: 396 is the determined cDNA sequence for clone 26519.
 SEQ ID NO: 397 is the determined cDNA sequence for clone 26520.
 SEQ ID NO: 398 is the determined cDNA sequence for clone 26521.
 SEQ ID NO: 399 is the determined cDNA sequence for clone 26522.
 15 SEQ ID NO: 400 is the determined cDNA sequence for clone 26523.
 SEQ ID NO: 401 is the determined cDNA sequence for clone 26524.
 SEQ ID NO: 402 is the determined cDNA sequence for clone 26526.
 SEQ ID NO: 403 is the determined cDNA sequence for clone 26527.
 SEQ ID NO: 404 is the determined cDNA sequence for clone 26528.
 20 SEQ ID NO: 405 is the determined cDNA sequence for clone 26529.
 SEQ ID NO: 406 is the determined cDNA sequence for clone 26530.
 SEQ ID NO: 407 is the determined cDNA sequence for clone 26532.
 SEQ ID NO: 408 is the determined cDNA sequence for clone 26533.
 SEQ ID NO: 409 is the determined cDNA sequence for clone 26534.
 25 SEQ ID NO: 410 is the determined cDNA sequence for clone 26535.
 SEQ ID NO: 411 is the determined cDNA sequence for clone 26536.
 SEQ ID NO: 412 is the determined cDNA sequence for clone 26537.
 SEQ ID NO: 413 is the determined cDNA sequence for clone 26538.
 SEQ ID NO: 414 is the determined cDNA sequence for clone 26540.

SEQ ID NO: 415 is the determined cDNA sequence for clone 26541.
 SEQ ID NO: 416 is the determined cDNA sequence for clone 26542.
 SEQ ID NO: 417 is the determined cDNA sequence for clone 26543.
 SEQ ID NO: 418 is the determined cDNA sequence for clone 26544.
 5 SEQ ID NO: 419 is the determined cDNA sequence for clone 26546.
 SEQ ID NO: 420 is the determined cDNA sequence for clone 26547.
 SEQ ID NO: 421 is the determined cDNA sequence for clone 26548.
 SEQ ID NO: 422 is the determined cDNA sequence for clone 26549.
 SEQ ID NO: 423 is the determined cDNA sequence for clone 26550.
 10 SEQ ID NO: 424 is the determined cDNA sequence for clone 26551.
 SEQ ID NO: 425 is the determined cDNA sequence for clone 26552.
 SEQ ID NO: 426 is the determined cDNA sequence for clone 26553.
 SEQ ID NO: 427 is the determined cDNA sequence for clone 26554.
 SEQ ID NO: 428 is the determined cDNA sequence for clone 26556.
 15 SEQ ID NO: 429 is the determined cDNA sequence for clone 26557.
 SEQ ID NO: 430 is the determined cDNA sequence for clone 27631.
 SEQ ID NO: 431 is the determined cDNA sequence for clone 27632.
 SEQ ID NO: 432 is the determined cDNA sequence for clone 27633.
 SEQ ID NO: 433 is the determined cDNA sequence for clone 27635.
 20 SEQ ID NO: 434 is the determined cDNA sequence for clone 27636.
 SEQ ID NO: 435 is the determined cDNA sequence for clone 27637.
 SEQ ID NO: 436 is the determined cDNA sequence for clone 27638.
 SEQ ID NO: 437 is the determined cDNA sequence for clone 27639.
 SEQ ID NO: 438 is the determined cDNA sequence for clone 27640.
 25 SEQ ID NO: 439 is the determined cDNA sequence for clone 27641.
 SEQ ID NO: 440 is the determined cDNA sequence for clone 27642.
 SEQ ID NO: 441 is the determined cDNA sequence for clone 27644.
 SEQ ID NO: 442 is the determined cDNA sequence for clone 27646.
 SEQ ID NO: 443 is the determined cDNA sequence for clone 27647.

SEQ ID NO: 444 is the determined cDNA sequence for clone 27649.
 SEQ ID NO: 445 is the determined cDNA sequence for clone 27650.
 SEQ ID NO: 446 is the determined cDNA sequence for clone 27651.
 SEQ ID NO: 447 is the determined cDNA sequence for clone 27652.
 5 SEQ ID NO: 448 is the determined cDNA sequence for clone 27654.
 SEQ ID NO: 449 is the determined cDNA sequence for clone 27655.
 SEQ ID NO: 450 is the determined cDNA sequence for clone 27657.
 SEQ ID NO: 451 is the determined cDNA sequence for clone 27659.
 SEQ ID NO: 452 is the determined cDNA sequence for clone 27665.
 10 SEQ ID NO: 453 is the determined cDNA sequence for clone 27666.
 SEQ ID NO: 454 is the determined cDNA sequence for clone 27668.
 SEQ ID NO: 455 is the determined cDNA sequence for clone 27670.
 SEQ ID NO: 456 is the determined cDNA sequence for clone 27671.
 SEQ ID NO: 457 is the determined cDNA sequence for clone 27672.
 15 SEQ ID NO: 458 is the determined cDNA sequence for clone 27674.
 SEQ ID NO: 459 is the determined cDNA sequence for clone 27677.
 SEQ ID NO: 460 is the determined cDNA sequence for clone 27681.
 SEQ ID NO: 461 is the determined cDNA sequence for clone 27682.
 SEQ ID NO: 462 is the determined cDNA sequence for clone 27683.
 20 SEQ ID NO: 463 is the determined cDNA sequence for clone 27686.
 SEQ ID NO: 464 is the determined cDNA sequence for clone 27688.
 SEQ ID NO: 465 is the determined cDNA sequence for clone 27689.
 SEQ ID NO: 466 is the determined cDNA sequence for clone 27690.
 SEQ ID NO: 467 is the determined cDNA sequence for clone 27693.
 25 SEQ ID NO: 468 is the determined cDNA sequence for clone 27699.
 SEQ ID NO: 469 is the determined cDNA sequence for clone 27700.
 SEQ ID NO: 470 is the determined cDNA sequence for clone 27702.
 SEQ ID NO: 471 is the determined cDNA sequence for clone 27705.
 SEQ ID NO: 472 is the determined cDNA sequence for clone 27706.

SEQ ID NO: 473 is the determined cDNA sequence for clone 27707.
 SEQ ID NO: 474 is the determined cDNA sequence for clone 27708.
 SEQ ID NO: 475 is the determined cDNA sequence for clone 27709.
 SEQ ID NO: 476 is the determined cDNA sequence for clone 27710.
 5 SEQ ID NO: 477 is the determined cDNA sequence for clone 27711.
 SEQ ID NO: 478 is the determined cDNA sequence for clone 27712.
 SEQ ID NO: 479 is the determined cDNA sequence for clone 27713.
 SEQ ID NO: 480 is the determined cDNA sequence for clone 27714.
 SEQ ID NO: 481 is the determined cDNA sequence for clone 27715.
 10 SEQ ID NO: 482 is the determined cDNA sequence for clone 27716.
 SEQ ID NO: 483 is the determined cDNA sequence for clone 27717.
 SEQ ID NO: 484 is the determined cDNA sequence for clone 27718.
 SEQ ID NO: 485 is the determined cDNA sequence for clone 27719.
 SEQ ID NO: 486 is the determined cDNA sequence for clone 27720.
 15 SEQ ID NO: 487 is the determined cDNA sequence for clone 27722.
 SEQ ID NO: 488 is the determined cDNA sequence for clone 27723.
 SEQ ID NO: 489 is the determined cDNA sequence for clone 27724.
 SEQ ID NO: 490 is the determined cDNA sequence for clone 27726.
 SEQ ID NO: 491 is the determined cDNA sequence for clone 25015.
 20 SEQ ID NO: 492 is the determined cDNA sequence for clone 25016.
 SEQ ID NO: 493 is the determined cDNA sequence for clone 25017.
 SEQ ID NO: 494 is the determined cDNA sequence for clone 25018
 SEQ ID NO: 495 is the determined cDNA sequence for clone 25030.
 SEQ ID NO: 496 is the determined cDNA sequence for clone 25033.
 25 SEQ ID NO: 497 is the determined cDNA sequence for clone 25034.
 SEQ ID NO: 498 is the determined cDNA sequence for clone 25035.
 SEQ ID NO: 499 is the determined cDNA sequence for clone 25036.
 SEQ ID NO: 500 is the determined cDNA sequence for clone 25037.
 SEQ ID NO: 501 is the determined cDNA sequence for clone 25038.

SEQ ID NO: 502 is the determined cDNA sequence for clone 25039.
 SEQ ID NO: 503 is the determined cDNA sequence for clone 25040.
 SEQ ID NO: 504 is the determined cDNA sequence for clone 25042.
 SEQ ID NO: 505 is the determined cDNA sequence for clone 25043.
 5 SEQ ID NO: 506 is the determined cDNA sequence for clone 25044.
 SEQ ID NO: 507 is the determined cDNA sequence for clone 25045.
 SEQ ID NO: 508 is the determined cDNA sequence for clone 25047.
 SEQ ID NO: 509 is the determined cDNA sequence for clone 25048.
 SEQ ID NO: 510 is the determined cDNA sequence for clone 25049.
 10 SEQ ID NO: 511 is the determined cDNA sequence for clone 25185.
 SEQ ID NO: 512 is the determined cDNA sequence for clone 25186.
 SEQ ID NO: 513 is the determined cDNA sequence for clone 25187.
 SEQ ID NO: 514 is the determined cDNA sequence for clone 25188.
 SEQ ID NO: 515 is the determined cDNA sequence for clone 25189.
 15 SEQ ID NO: 516 is the determined cDNA sequence for clone 25190.
 SEQ ID NO: 517 is the determined cDNA sequence for clone 25193.
 SEQ ID NO: 518 is the determined cDNA sequence for clone 25194.
 SEQ ID NO: 519 is the determined cDNA sequence for clone 25196.
 SEQ ID NO: 520 is the determined cDNA sequence for clone 25198.
 20 SEQ ID NO: 521 is the determined cDNA sequence for clone 25199.
 SEQ ID NO: 522 is the determined cDNA sequence for clone 25200.
 SEQ ID NO: 523 is the determined cDNA sequence for clone 25202.
 SEQ ID NO: 524 is the determined cDNA sequence for clone 25364.
 SEQ ID NO: 525 is the determined cDNA sequence for clone 25366.
 25 SEQ ID NO: 526 is the determined cDNA sequence for clone 25367.
 SEQ ID NO: 527 is the determined cDNA sequence for clone 25368.
 SEQ ID NO: 528 is the determined cDNA sequence for clone 25369.
 SEQ ID NO: 529 is the determined cDNA sequence for clone 25370.
 SEQ ID NO: 530 is the determined cDNA sequence for clone 25371.

SEQ ID NO: 531 is the determined cDNA sequence for clone 25372.
 SEQ ID NO: 532 is the determined cDNA sequence for clone 25373.
 SEQ ID NO: 533 is the determined cDNA sequence for clone 25374.
 SEQ ID NO: 534 is the determined cDNA sequence for clone 25376.
 5 SEQ ID NO: 535 is the determined cDNA sequence for clone 25377.
 SEQ ID NO: 536 is the determined cDNA sequence for clone 25378.
 SEQ ID NO: 537 is the determined cDNA sequence for clone 25379.
 SEQ ID NO: 538 is the determined cDNA sequence for clone 25380.
 SEQ ID NO: 539 is the determined cDNA sequence for clone 25381.
 10 SEQ ID NO: 540 is the determined cDNA sequence for clone 25382.
 SEQ ID NO: 541 is the determined cDNA sequence for clone 25383.
 SEQ ID NO: 542 is the determined cDNA sequence for clone 25385.
 SEQ ID NO: 543 is the determined cDNA sequence for clone 25386.
 SEQ ID NO: 544 is the determined cDNA sequence for clone 25387.
 15 SEQ ID NO: 545 is the determined cDNA sequence for clone 26013.
 SEQ ID NO: 546 is the determined cDNA sequence for clone 26014.
 SEQ ID NO: 547 is the determined cDNA sequence for clone 26016.
 SEQ ID NO: 548 is the determined cDNA sequence for clone 26017.
 SEQ ID NO: 549 is the determined cDNA sequence for clone 26018.
 20 SEQ ID NO: 550 is the determined cDNA sequence for clone 26019.
 SEQ ID NO: 551 is the determined cDNA sequence for clone 26020.
 SEQ ID NO: 552 is the determined cDNA sequence for clone 26021.
 SEQ ID NO: 553 is the determined cDNA sequence for clone 26022.
 SEQ ID NO: 554 is the determined cDNA sequence for clone 26027.
 25 SEQ ID NO: 555 is the determined cDNA sequence for clone 26197.
 SEQ ID NO: 556 is the determined cDNA sequence for clone 26199.
 SEQ ID NO: 557 is the determined cDNA sequence for clone 26201.
 SEQ ID NO: 558 is the determined cDNA sequence for clone 26202.
 SEQ ID NO: 559 is the determined cDNA sequence for clone 26203.

SEQ ID NO: 560 is the determined cDNA sequence for clone 26204.
 SEQ ID NO: 561 is the determined cDNA sequence for clone 26205.
 SEQ ID NO: 562 is the determined cDNA sequence for clone 26206.
 SEQ ID NO: 563 is the determined cDNA sequence for clone 26208.
 5 SEQ ID NO: 564 is the determined cDNA sequence for clone 26211.
 SEQ ID NO: 565 is the determined cDNA sequence for clone 26212.
 SEQ ID NO: 566 is the determined cDNA sequence for clone 26213.
 SEQ ID NO: 567 is the determined cDNA sequence for clone 26214.
 SEQ ID NO: 568 is the determined cDNA sequence for clone 26215.
 10 SEQ ID NO: 569 is the determined cDNA sequence for clone 26216.
 SEQ ID NO: 570 is the determined cDNA sequence for clone 26217.
 SEQ ID NO: 571 is the determined cDNA sequence for clone 26218.
 SEQ ID NO: 572 is the determined cDNA sequence for clone 26219.
 SEQ ID NO: 573 is the determined cDNA sequence for clone 26220.
 15 SEQ ID NO: 574 is the determined cDNA sequence for clone 26221.
 SEQ ID NO: 575 is the determined cDNA sequence for clone 26224.
 SEQ ID NO: 576 is the determined cDNA sequence for clone 26225.
 SEQ ID NO: 577 is the determined cDNA sequence for clone 26226.
 SEQ ID NO: 578 is the determined cDNA sequence for clone 26227.
 20 SEQ ID NO: 579 is the determined cDNA sequence for clone 26228.
 SEQ ID NO: 580 is the determined cDNA sequence for clone 26230.
 SEQ ID NO: 581 is the determined cDNA sequence for clone 26231.
 SEQ ID NO: 582 is the determined cDNA sequence for clone 26234.
 SEQ ID NO: 583 is the determined cDNA sequence for clone 26236.
 25 SEQ ID NO: 584 is the determined cDNA sequence for clone 26237.
 SEQ ID NO: 585 is the determined cDNA sequence for clone 26239.
 SEQ ID NO: 586 is the determined cDNA sequence for clone 26240.
 SEQ ID NO: 587 is the determined cDNA sequence for clone 26241.
 SEQ ID NO: 588 is the determined cDNA sequence for clone 26242.

SEQ ID NO: 589 is the determined cDNA sequence for clone 26246.
 SEQ ID NO: 590 is the determined cDNA sequence for clone 26247.
 SEQ ID NO: 591 is the determined cDNA sequence for clone 26248.
 SEQ ID NO: 592 is the determined cDNA sequence for clone 26249.
 5 SEQ ID NO: 593 is the determined cDNA sequence for clone 26250.
 SEQ ID NO: 594 is the determined cDNA sequence for clone 26251.
 SEQ ID NO: 595 is the determined cDNA sequence for clone 26252.
 SEQ ID NO: 596 is the determined cDNA sequence for clone 26253.
 SEQ ID NO: 597 is the determined cDNA sequence for clone 26254.
 10 SEQ ID NO: 598 is the determined cDNA sequence for clone 26255.
 SEQ ID NO: 599 is the determined cDNA sequence for clone 26256.
 SEQ ID NO: 600 is the determined cDNA sequence for clone 26257.
 SEQ ID NO: 601 is the determined cDNA sequence for clone 26259.
 SEQ ID NO: 602 is the determined cDNA sequence for clone 26260.
 15 SEQ ID NO: 603 is the determined cDNA sequence for clone 26261.
 SEQ ID NO: 604 is the determined cDNA sequence for clone 26262.
 SEQ ID NO: 605 is the determined cDNA sequence for clone 26263.
 SEQ ID NO: 606 is the determined cDNA sequence for clone 26264.
 SEQ ID NO: 607 is the determined cDNA sequence for clone 26265.
 20 SEQ ID NO: 608 is the determined cDNA sequence for clone 26266.
 SEQ ID NO: 609 is the determined cDNA sequence for clone 26268.
 SEQ ID NO: 610 is the determined cDNA sequence for clone 26269.
 SEQ ID NO: 611 is the determined cDNA sequence for clone 26271.
 SEQ ID NO: 612 is the determined cDNA sequence for clone 26273.
 25 SEQ ID NO: 613 is the determined cDNA sequence for clone 26810.
 SEQ ID NO: 614 is the determined cDNA sequence for clone 26811.
 SEQ ID NO: 615 is the determined cDNA sequence for clone 26812.1.
 SEQ ID NO: 616 is the determined cDNA sequence for clone 26812.2.
 SEQ ID NO: 617 is the determined cDNA sequence for clone 26813.

SEQ ID NO: 618 is the determined cDNA sequence for clone 26814.
 SEQ ID NO: 619 is the determined cDNA sequence for clone 26815.
 SEQ ID NO: 620 is the determined cDNA sequence for clone 26816.
 SEQ ID NO: 621 is the determined cDNA sequence for clone 26818.
 5 SEQ ID NO: 622 is the determined cDNA sequence for clone 26819.
 SEQ ID NO: 623 is the determined cDNA sequence for clone 26820.
 SEQ ID NO: 624 is the determined cDNA sequence for clone 26821.
 SEQ ID NO: 625 is the determined cDNA sequence for clone 26822.
 SEQ ID NO: 626 is the determined cDNA sequence for clone 26824.
 10 SEQ ID NO: 627 is the determined cDNA sequence for clone 26825.
 SEQ ID NO: 628 is the determined cDNA sequence for clone 26826.
 SEQ ID NO: 629 is the determined cDNA sequence for clone 26827.
 SEQ ID NO: 630 is the determined cDNA sequence for clone 26829.
 SEQ ID NO: 631 is the determined cDNA sequence for clone 26830.
 15 SEQ ID NO: 632 is the determined cDNA sequence for clone 26831.
 SEQ ID NO: 633 is the determined cDNA sequence for clone 26832.
 SEQ ID NO: 634 is the determined cDNA sequence for clone 26835.
 SEQ ID NO: 635 is the determined cDNA sequence for clone 26836.
 SEQ ID NO: 636 is the determined cDNA sequence for clone 26837.
 20 SEQ ID NO: 637 is the determined cDNA sequence for clone 26839.
 SEQ ID NO: 638 is the determined cDNA sequence for clone 26841.
 SEQ ID NO: 639 is the determined cDNA sequence for clone 26843.
 SEQ ID NO: 640 is the determined cDNA sequence for clone 26844.
 SEQ ID NO: 641 is the determined cDNA sequence for clone 26845.
 25 SEQ ID NO: 642 is the determined cDNA sequence for clone 26846.
 SEQ ID NO: 643 is the determined cDNA sequence for clone 26847.
 SEQ ID NO: 644 is the determined cDNA sequence for clone 26848.
 SEQ ID NO: 645 is the determined cDNA sequence for clone 26849.
 SEQ ID NO: 646 is the determined cDNA sequence for clone 26850.

SEQ ID NO: 647 is the determined cDNA sequence for clone 26851.
 SEQ ID NO: 648 is the determined cDNA sequence for clone 26852.
 SEQ ID NO: 649 is the determined cDNA sequence for clone 26853.
 SEQ ID NO: 650 is the determined cDNA sequence for clone 26854.
 5 SEQ ID NO: 651 is the determined cDNA sequence for clone 26856.
 SEQ ID NO: 652 is the determined cDNA sequence for clone 26857.
 SEQ ID NO: 653 is the determined cDNA sequence for clone 26858.
 SEQ ID NO: 654 is the determined cDNA sequence for clone 26859.
 SEQ ID NO: 655 is the determined cDNA sequence for clone 26860.
 10 SEQ ID NO: 656 is the determined cDNA sequence for clone 26862.
 SEQ ID NO: 657 is the determined cDNA sequence for clone 26863.
 SEQ ID NO: 658 is the determined cDNA sequence for clone 26864.
 SEQ ID NO: 659 is the determined cDNA sequence for clone 26865.
 SEQ ID NO: 660 is the determined cDNA sequence for clone 26867.
 15 SEQ ID NO: 661 is the determined cDNA sequence for clone 26868.
 SEQ ID NO: 662 is the determined cDNA sequence for clone 26871.
 SEQ ID NO: 663 is the determined cDNA sequence for clone 26873.
 SEQ ID NO: 664 is the determined cDNA sequence for clone 26875.
 SEQ ID NO: 665 is the determined cDNA sequence for clone 26876.
 20 SEQ ID NO: 666 is the determined cDNA sequence for clone 26877.
 SEQ ID NO: 667 is the determined cDNA sequence for clone 26878.
 SEQ ID NO: 668 is the determined cDNA sequence for clone 26880.
 SEQ ID NO: 669 is the determined cDNA sequence for clone 26882.
 SEQ ID NO: 670 is the determined cDNA sequence for clone 26883.
 25 SEQ ID NO: 671 is the determined cDNA sequence for clone 26884.
 SEQ ID NO: 672 is the determined cDNA sequence for clone 26885.
 SEQ ID NO: 673 is the determined cDNA sequence for clone 26886.
 SEQ ID NO: 674 is the determined cDNA sequence for clone 26887.
 SEQ ID NO: 675 is the determined cDNA sequence for clone 26888.

SEQ ID NO: 676 is the determined cDNA sequence for clone 26889.
 SEQ ID NO: 677 is the determined cDNA sequence for clone 26890.
 SEQ ID NO: 678 is the determined cDNA sequence for clone 26892.
 SEQ ID NO: 679 is the determined cDNA sequence for clone 26894.
 5 SEQ ID NO: 680 is the determined cDNA sequence for clone 26895.
 SEQ ID NO: 681 is the determined cDNA sequence for clone 26897.
 SEQ ID NO: 682 is the determined cDNA sequence for clone 26898.
 SEQ ID NO: 683 is the determined cDNA sequence for clone 26899.
 SEQ ID NO: 684 is the determined cDNA sequence for clone 26900.
 10 SEQ ID NO: 685 is the determined cDNA sequence for clone 26901.
 SEQ ID NO: 686 is the determined cDNA sequence for clone 26903.
 SEQ ID NO: 687 is the determined cDNA sequence for clone 26905.
 SEQ ID NO: 688 is the determined cDNA sequence for clone 26906.
 SEQ ID NO: 689 is the determined cDNA sequence for clone 26708.
 15 SEQ ID NO: 690 is the determined cDNA sequence for clone 26709.
 SEQ ID NO: 691 is the determined cDNA sequence for clone 26710.
 SEQ ID NO: 692 is the determined cDNA sequence for clone 26711.
 SEQ ID NO: 693 is the determined cDNA sequence for clone 26712.
 SEQ ID NO: 694 is the determined cDNA sequence for clone 26713.
 20 SEQ ID NO: 695 is the determined cDNA sequence for clone 26714.
 SEQ ID NO: 696 is the determined cDNA sequence for clone 26715.
 SEQ ID NO: 697 is the determined cDNA sequence for clone 26716.
 SEQ ID NO: 698 is the determined cDNA sequence for clone 26717.
 SEQ ID NO: 699 is the determined cDNA sequence for clone 26718.
 25 SEQ ID NO: 700 is the determined cDNA sequence for clone 26719.
 SEQ ID NO: 701 is the determined cDNA sequence for clone 26720.
 SEQ ID NO: 702 is the determined cDNA sequence for clone 26721.
 SEQ ID NO: 703 is the determined cDNA sequence for clone 26722.
 SEQ ID NO: 704 is the determined cDNA sequence for clone 26723.

SEQ ID NO: 705 is the determined cDNA sequence for clone 26724.
 SEQ ID NO: 706 is the determined cDNA sequence for clone 26725.
 SEQ ID NO: 707 is the determined cDNA sequence for clone 26726.
 SEQ ID NO: 708 is the determined cDNA sequence for clone 26727.
 5 SEQ ID NO: 709 is the determined cDNA sequence for clone 26728.
 SEQ ID NO: 710 is the determined cDNA sequence for clone 26729.
 SEQ ID NO: 711 is the determined cDNA sequence for clone 26730.
 SEQ ID NO: 712 is the determined cDNA sequence for clone 26731.
 SEQ ID NO: 713 is the determined cDNA sequence for clone 26732.
 10 SEQ ID NO: 714 is the determined cDNA sequence for clone 26733.1.
 SEQ ID NO: 715 is the determined cDNA sequence for clone 26733.2.
 SEQ ID NO: 716 is the determined cDNA sequence for clone 26734.
 SEQ ID NO: 717 is the determined cDNA sequence for clone 26735.
 SEQ ID NO: 718 is the determined cDNA sequence for clone 26736.
 15 SEQ ID NO: 719 is the determined cDNA sequence for clone 26737.
 SEQ ID NO: 720 is the determined cDNA sequence for clone 26738.
 SEQ ID NO: 721 is the determined cDNA sequence for clone 26739.
 SEQ ID NO: 722 is the determined cDNA sequence for clone 26741.
 SEQ ID NO: 723 is the determined cDNA sequence for clone 26742.
 20 SEQ ID NO: 724 is the determined cDNA sequence for clone 26743.
 SEQ ID NO: 725 is the determined cDNA sequence for clone 26744.
 SEQ ID NO: 726 is the determined cDNA sequence for clone 26745.
 SEQ ID NO: 727 is the determined cDNA sequence for clone 26746.
 SEQ ID NO: 728 is the determined cDNA sequence for clone 26747.
 25 SEQ ID NO: 729 is the determined cDNA sequence for clone 26748.
 SEQ ID NO: 730 is the determined cDNA sequence for clone 26749.
 SEQ ID NO: 731 is the determined cDNA sequence for clone 26750.
 SEQ ID NO: 732 is the determined cDNA sequence for clone 26751.
 SEQ ID NO: 733 is the determined cDNA sequence for clone 26752.

SEQ ID NO: 734 is the determined cDNA sequence for clone 26753.
 SEQ ID NO: 735 is the determined cDNA sequence for clone 26754.
 SEQ ID NO: 736 is the determined cDNA sequence for clone 26755.
 SEQ ID NO: 737 is the determined cDNA sequence for clone 26756.
 5 SEQ ID NO: 738 is the determined cDNA sequence for clone 26757.
 SEQ ID NO: 739 is the determined cDNA sequence for clone 26758.
 SEQ ID NO: 740 is the determined cDNA sequence for clone 26759.
 SEQ ID NO: 741 is the determined cDNA sequence for clone 26760.
 SEQ ID NO: 742 is the determined cDNA sequence for clone 26761.
 10 SEQ ID NO: 743 is the determined cDNA sequence for clone 26762.
 SEQ ID NO: 744 is the determined cDNA sequence for clone 26763.
 SEQ ID NO: 745 is the determined cDNA sequence for clone 26764.
 SEQ ID NO: 746 is the determined cDNA sequence for clone 26765.
 SEQ ID NO: 747 is the determined cDNA sequence for clone 26766.
 15 SEQ ID NO: 748 is the determined cDNA sequence for clone 26767.
 SEQ ID NO: 749 is the determined cDNA sequence for clone 26768.
 SEQ ID NO: 750 is the determined cDNA sequence for clone 26769.
 SEQ ID NO: 751 is the determined cDNA sequence for clone 26770.
 SEQ ID NO: 752 is the determined cDNA sequence for clone 26771.
 20 SEQ ID NO: 753 is the determined cDNA sequence for clone 26772.
 SEQ ID NO: 754 is the determined cDNA sequence for clone 26773.
 SEQ ID NO: 755 is the determined cDNA sequence for clone 26774.
 SEQ ID NO: 756 is the determined cDNA sequence for clone 26775.
 SEQ ID NO: 757 is the determined cDNA sequence for clone 26776.
 25 SEQ ID NO: 758 is the determined cDNA sequence for clone 26777.
 SEQ ID NO: 759 is the determined cDNA sequence for clone 26778.
 SEQ ID NO: 760 is the determined cDNA sequence for clone 26779.
 SEQ ID NO: 761 is the determined cDNA sequence for clone 26781.
 SEQ ID NO: 762 is the determined cDNA sequence for clone 26782.

SEQ ID NO: 763 is the determined cDNA sequence for clone 26783.
 SEQ ID NO: 764 is the determined cDNA sequence for clone 26784.
 SEQ ID NO: 765 is the determined cDNA sequence for clone 26785.
 SEQ ID NO: 766 is the determined cDNA sequence for clone 26786.
 5 SEQ ID NO: 767 is the determined cDNA sequence for clone 26787.
 SEQ ID NO: 768 is the determined cDNA sequence for clone 26788.
 SEQ ID NO: 769 is the determined cDNA sequence for clone 26790.
 SEQ ID NO: 770 is the determined cDNA sequence for clone 26791.
 SEQ ID NO: 771 is the determined cDNA sequence for clone 26792.
 10 SEQ ID NO: 772 is the determined cDNA sequence for clone 26793.
 SEQ ID NO: 773 is the determined cDNA sequence for clone 26794.
 SEQ ID NO: 774 is the determined cDNA sequence for clone 26795.
 SEQ ID NO: 775 is the determined cDNA sequence for clone 26796.
 SEQ ID NO: 776 is the determined cDNA sequence for clone 26797.
 15 SEQ ID NO: 777 is the determined cDNA sequence for clone 26798.
 SEQ ID NO: 778 is the determined cDNA sequence for clone 26800.
 SEQ ID NO: 779 is the determined cDNA sequence for clone 26801.
 SEQ ID NO: 780 is the determined cDNA sequence for clone 26802.
 SEQ ID NO: 781 is the determined cDNA sequence for clone 26803.
 20 SEQ ID NO: 782 is the determined cDNA sequence for clone 26804.
 SEQ ID NO: 783 is the amino acid sequence for L773P.
 SEQ ID NO: 784 is the determined DNA sequence of the L773P expression
 construct.
 SEQ ID NO: 785 is the determined DNA sequence of the L773PA
 25 expression construct.
 SEQ ID NO: 786 is a predicted amino acid sequence for L552S.
 SEQ ID NO: 787 is a predicted amino acid sequence for L840P.
 SEQ ID NO: 788 is the full-length cDNA sequence for L548S.
 SEQ ID NO: 789 is the amino acid sequence encoded by SEQ ID NO: 788.

SEQ ID NO: 791 is the predicted amino acid sequence encoded by the cDNA sequence of SEQ ID NO: 790.

SEQ ID NO: 792 is the determined cDNA sequence for an isoform of L552S.

SEQ ID NO: 793 is the predicted amino acid sequence encoded by SEQ ID NO: 792.

SEQ ID NO: 794 is an extended cDNA sequence for L840P.

SEQ ID NO: 795 is the predicted amino acid sequence encoded by SEQ ID NO: 794.

SEQ ID NO: 796 is an extended cDNA sequence for L801P.

SEQ ID NO: 797 is a first predicted amino acid sequence encoded by SEQ ID NO: 796.

SEQ ID NO: 798 is a second predicted amino acid sequence encoded by SEQ ID NO: 796.

SEQ ID NO: 799 is a third predicted amino acid sequence encoded by SEQ ID NO: 796.

SEQ ID NO: 800 is the determined full-length sequence for L844P.

SEQ ID NO: 801 is the 5' consensus cDNA sequence for L551S.

SEQ ID NO: 802 is the 3' consensus cDNA sequence for L551S.

SEQ ID NO: 803 is the cDNA sequence for STY8.

SEQ ID NO: 804 is an extended cDNA sequence for L551S.

SEQ ID NO: 805 is the amino acid sequence for STY8.

SEQ ID NO: 806 is the extended amino acid sequence for L551S.

SEQ ID NO: 807 is the determined full-length cDNA sequence for L773P.

SEQ ID NO: 808 is the full-length cDNA sequence of L552S.

SEQ ID NO: 809 is the full-length amino acid sequence of L552S.

SEQ ID NO: 810 is the determined cDNA sequence of clone 50989.

SEQ ID NO: 811 is the determined cDNA sequence of clone 50990.

SEQ ID NO: 812 is the determined cDNA sequence of clone 50992.

SEQ ID NO: 813-824 are the determined cDNA sequences for clones isolated from lung tumor tissue.

SEQ ID NO: 825 is the determined cDNA sequence for the full-length
5 L551S clone 54305.

SEQ ID NO: 826 is the determined cDNA sequence for the full-length
L551S clone 54298.

SEQ ID NO: 827 is the full-length amino acid sequence for L551S.

Tables 1-6 contain the sequence identifiers for SEQ ID NO:878-1664.

Table 1A

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 828 | R0126:A02 | 869 | R0126:D12 |
| 829 | R0126:A03 | 870 | R0126:E01 |
| 830 | R0126:A05 | 871 | R0126:E02 |
| 831 | R0126:A06 | 872 | R0126:E03 |
| 832 | R0126:A08 | 873 | R0126:E04 |
| 833 | R0126:A09 | 874 | R0126:E05 |
| 834 | R0126:A10 | 875 | R0126:E06 |
| 835 | R0126:A11 | 876 | R0126:E07 |
| 836 | R0126:A12 | 877 | R0126:E08 |
| 837 | R0126:B01 | 878 | R0126:E09 |
| 838 | R0126:B03 | 879 | R0126:E10 |
| 839 | R0126:B04 | 880 | R0126:E11 |
| 840 | R0126:B05 | 881 | R0126:E12 |
| 841 | R0126:B06 | 882 | R0126:F01 |
| 842 | R0126:B07 | 883 | R0126:F02 |
| 843 | R0126:B08 | 884 | R0126:F03 |
| 844 | R0126:B09 | 885 | R0126:F04 |
| 845 | R0126:B11 | 886 | R0126:F05 |
| 846 | R0126:B12 | 887 | R0126:F06 |
| 847 | R0126:C01 | 888 | R0126:F07 |
| 848 | R0126:C02 | 889 | R0126:F08 |
| 849 | R0126:C03 | 890 | R0126:F10 |
| 850 | R0126:C05 | 891 | R0126:F11 |
| 851 | R0126:C06 | 892 | R0126:F12 |
| 852 | R0126:C07 | 893 | R0126:G01 |
| 853 | R0126:C08 | 894 | R0126:G02 |
| 854 | R0126:C09 | 895 | R0126:G03 |
| 855 | R0126:C10 | 896 | R0126:G04 |
| 856 | R0126:C11 | 897 | R0126:G05 |
| 857 | R0126:C12 | 898 | R0126:G06 |
| 858 | R0126:D01 | 899 | R0126:G07 |
| 859 | R0126:D02 | 900 | R0126:G09 |
| 860 | R0126:D03 | 901 | R0126:G10 |
| 861 | R0126:D04 | 902 | R0126:G11 |
| 862 | R0126:D05 | 903 | R0126:G12 |
| 863 | R0126:D06 | 904 | R0126:H01 |
| 864 | R0126:D07 | 905 | R0126:H02 |
| 865 | R0126:D08 | 906 | R0126:H03 |
| 866 | R0126:D09 | 907 | R0126:H04 |

Table 1B

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 910 | R0126:H07 | 951 | R0127:D10 |
| 911 | R0126:H09 | 952 | R0127:D11 |
| 912 | R0126:H10 | 953 | R0127:D12 |
| 913 | R0126:H11 | 954 | R0127:E02 |
| 914 | R0127:A02 | 955 | R0127:E03 |
| 915 | R0127:A05 | 956 | R0127:E04 |
| 916 | R0127:A06 | 957 | R0127:E05 |
| 917 | R0127:A07 | 958 | R0127:E06 |
| 918 | R0127:A08 | 959 | R0127:E07 |
| 919 | R0127:A09 | 960 | R0127:E08 |
| 920 | R0127:A10 | 961 | R0127:E09 |
| 921 | R0127:A11 | 962 | R0127:E10 |
| 922 | R0127:A12 | 963 | R0127:E11 |
| 923 | R0127:B01 | 964 | R0127:F01 |
| 924 | R0127:B03 | 965 | R0127:F02 |
| 925 | R0127:B04 | 966 | R0127:F03 |
| 926 | R0127:B05 | 967 | R0127:F04 |
| 927 | R0127:B06 | 968 | R0127:F05 |
| 928 | R0127:B07 | 969 | R0127:F06 |
| 929 | R0127:B08 | 970 | R0127:F07 |
| 930 | R0127:B09 | 971 | R0127:F08 |
| 931 | R0127:B10 | 972 | R0127:F10 |
| 932 | R0127:B11 | 973 | R0127:F11 |
| 933 | R0127:B12 | 974 | R0127:F12 |
| 934 | R0127:C01 | 975 | R0127:G01 |
| 935 | R0127:C03 | 976 | R0127:G02 |
| 936 | R0127:C04 | 977 | R0127:G03 |
| 937 | R0127:C05 | 978 | R0127:G04 |
| 938 | R0127:C07 | 979 | R0127:G05 |
| 939 | R0127:C08 | 980 | R0127:G06 |
| 940 | R0127:C09 | 981 | R0127:G07 |
| 941 | R0127:C10 | 982 | R0127:G08 |
| 942 | R0127:C11 | 983 | R0127:G09 |
| 943 | R0127:D01 | 984 | R0127:G10 |
| 944 | R0127:D02 | 985 | R0127:G11 |
| 945 | R0127:D03 | 986 | R0127:G12 |
| 946 | R0127:D04 | 987 | R0127:H01 |
| 947 | R0127:D05 | 988 | R0127:H02 |
| 948 | R0127:D06 | 989 | R0127:H03 |

Table 1C

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 992 | R0127:H06 | 1034 | R0128:D11 |
| 993 | R0127:H07 | 1035 | R0128:D12 |
| 994 | R0127:H08 | 1036 | R0128:E01 |
| 995 | R1027:H09 | 1037 | R0128:E02 |
| 996 | R1027:H10 | 1038 | R0128:E03 |
| 997 | R1027:H11 | 1039 | R0128:E04 |
| 998 | R1028:A02 | 1040 | R0128:E05 |
| 999 | R1028:A05 | 1041 | R0128:E06 |
| 1000 | R1028:A06 | 1042 | R0128:E07 |
| 1001 | R1028:A07 | 1043 | R0128:E08 |
| 1002 | R1028:A08 | 1044 | R0128:E09 |
| 1003 | R1028:A09 | 1045 | R0128:E10 |
| 1004 | R1028:A10 | 1046 | R0128:E12 |
| 1005 | R1028:B01 | 1047 | R0128:F01 |
| 1006 | R1028:B02 | 1048 | R0128:F02 |
| 1007 | R1028:B03 | 1049 | R0128:F03 |
| 1008 | R1028:B04 | 1050 | R0128:F04 |
| 1009 | R1028:B05 | 1051 | R0128:F06 |
| 1010 | R1028:B08 | 1052 | R0128:F07 |
| 1011 | R1028:B09 | 1053 | R0128:F08 |
| 1012 | R1028:B10 | 1054 | R0128:F09 |
| 1013 | R1028:B11 | 1055 | R0128:F10 |
| 1014 | R1028:B12 | 1056 | R0128:F12 |
| 1015 | R1028:C01 | 1057 | R0128:G01 |
| 1016 | R1028:C03 | 1058 | R0128:G02 |
| 1017 | R1028:C04 | 1059 | R0128:G03 |
| 1018 | R1028:C05 | 1060 | R0128:G04 |
| 1019 | R1028:C06 | 1061 | R0128:G05 |
| 1020 | R1028:C07 | 1062 | R0128:G06 |
| 1021 | R1028:C08 | 1063 | R0128:G07 |
| 1022 | R1028:C10 | 1064 | R0128:G09 |
| 1023 | R1028:C11 | 1065 | R0128:G10 |
| 1024 | R1028:C12 | 1066 | R0128:G11 |
| 1025 | R1028:D01 | 1067 | R0128:G12 |
| 1026 | R1028:D02 | 1068 | R0128:H01 |
| 1027 | R1028:D04 | 1069 | R0128:H02 |
| 1028 | R1028:D05 | 1070 | R0128:H03 |
| 1029 | R1028:D06 | 1071 | R0128:H04 |
| 1030 | R1028:D07 | 1072 | R0128:H05 |
| 1031 | R1028:D08 | 1073 | R0128:H06 |

44

Table 1D

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1076 | R0128:H09 | 1118 | R0130:D12 |
| 1077 | R0128:H10 | 1119 | R0130:E01 |
| 1078 | R0128:H11 | 1120 | R0130:E02 |
| 1079 | R0130:A02 | 1121 | R0130:E03 |
| 1080 | R0130:A05 | 1122 | R0130:E04 |
| 1081 | R0130:A06 | 1123 | R0130:E05 |
| 1082 | R0130:A08 | 1124 | R0130:E06 |
| 1083 | R0130:A09 | 1125 | R0130:E07 |
| 1084 | R0130:A10 | 1126 | R0130:E08 |
| 1085 | R0130:A11 | 1127 | R0130:E09 |
| 1086 | R0130:A12 | 1128 | R0130:E10 |
| 1087 | R0130:B01 | 1129 | R0130:E11 |
| 1088 | R0130:B02 | 1130 | R0130:E12 |
| 1089 | R0130:B03 | 1131 | R0130:F02 |
| 1090 | R0130:B04 | 1132 | R0130:F03 |
| 1091 | R0130:B05 | 1133 | R0130:F05 |
| 1092 | R0130:B06 | 1134 | R0130:F06 |
| 1093 | R0130:B08 | 1135 | R0130:F07 |
| 1094 | R0130:B09 | 1136 | R0130:F08 |
| 1095 | R0130:B10 | 1137 | R0130:F09 |
| 1096 | R0130:B11 | 1138 | R0130:F10 |
| 1097 | R0130:B12 | 1139 | R0130:F11 |
| 1098 | R0130:C02 | 1140 | R0130:F12 |
| 1099 | R0130:C03 | 1141 | R0130:G01 |
| 1100 | R0130:C04 | 1142 | R0130:G02 |
| 1101 | R0130:C05 | 1143 | R0130:G03 |
| 1102 | R0130:C06 | 1144 | R0130:G04 |
| 1103 | R0130:C07 | 1145 | R0130:G05 |
| 1104 | R0130:C08 | 1146 | R0130:G06 |
| 1105 | R0130:C09 | 1147 | R0130:G07 |
| 1106 | R0130:C10 | 1148 | R0130:G08 |
| 1107 | R0130:C11 | 1149 | R0130:G09 |
| 1108 | R0130:C12 | 1150 | R0130:G10 |
| 1109 | R0130:D02 | 1151 | R0130:G11 |
| 1110 | R0130:D03 | 1152 | R0130:G12 |
| 1111 | R0130:D04 | 1153 | R0130:H01 |
| 1112 | R0130:D05 | 1154 | R0130:H02 |
| 1113 | R0130:D06 | 1155 | R0130:H04 |
| 1114 | R0130:D07 | 1156 | R0130:H05 |
| 1115 | R0130:D09 | 1157 | R0130:H06 |

[illegible]

Table 1E

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1160 | R0130:H09 | 1200 | R0131:E01 |
| 1161 | R0130:H10 | 1201 | R0131:E02 |
| 1162 | R0130:H11 | 1202 | R0131:E03 |
| 1163 | R0131:A02 | 1203 | R0131:E04 |
| 1164 | R0131:A05 | 1204 | R0131:E06 |
| 1165 | R0131:A06 | 1205 | R0131:E07 |
| 1166 | R0131:A07 | 1206 | R0131:E08 |
| 1167 | R0131:A08 | 1207 | R0131:E10 |
| 1168 | R0131:A09 | 1208 | R0131:E11 |
| 1169 | R0131:A11 | 1209 | R0131:E12 |
| 1170 | R0131:A12 | 1210 | R0131:F02 |
| 1171 | R0131:B02 | 1211 | R0131:F04 |
| 1172 | R0131:B03 | 1212 | R0131:F05 |
| 1173 | R0131:B04 | 1213 | R0131:F06 |
| 1174 | R0131:B05 | 1214 | R0131:F07 |
| 1175 | R0131:B07 | 1215 | R0131:F08 |
| 1176 | R0131:B08 | 1216 | R0131:F09 |
| 1177 | R0131:B09 | 1217 | R0131:F10 |
| 1178 | R0131:B10 | 1218 | R0131:F11 |
| 1179 | R0131:B11 | 1219 | R0131:F12 |
| 1180 | R0131:C01 | 1220 | R0131:G01 |
| 1181 | R0131:C02 | 1221 | R0131:G02 |
| 1182 | R0131:C03 | 1222 | R0131:G03 |
| 1183 | R0131:C04 | 1223 | R0131:G04 |
| 1184 | R0131:C06 | 1224 | R0131:G05 |
| 1185 | R0131:C07 | 1225 | R0131:G06 |
| 1186 | R0131:C08 | 1226 | R0131:G07 |
| 1187 | R0131:C10 | 1227 | R0131:G08 |
| 1188 | R0131:C11 | 1228 | R0131:G09 |
| 1189 | R0131:C12 | 1229 | R0131:G10 |
| 1190 | R0131:D02 | 1230 | R0131:G11 |
| 1191 | R0131:D03 | 1231 | R0131:G12 |
| 1192 | R0131:D04 | 1232 | R0131:H01 |
| 1193 | R0131:D05 | 1233 | R0131:H02 |
| 1194 | R0131:D06 | 1234 | R0131:H05 |
| 1195 | R0131:D07 | 1235 | R0131:H06 |
| 1196 | R0131:D09 | 1236 | R0131:H07 |
| 1197 | R0131:D10 | 1237 | R0131:H08 |
| 1198 | R0131:D11 | 1238 | R0131:H09 |
| 1199 | R0131:D12 | 1239 | R0131:H11 |

Table 2:
Clone names for NSCLC-SQL1 and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1240 | Contig 54 | | |
| 1241 | Contig 55 | | |
| 1242 | Contig 57 | | |
| 1243 | Contig 58 | | |
| 1244 | Contig 60 | | |
| 1245 | Contig 62 | | |
| 1246 | Contig 63 | | |
| 1247 | Contig 64 | | |
| 1248 | Contig 65 | | |
| 1249 | Contig 66 | | |
| 1250 | Contig 67 | | |
| 1251 | Contig 68 | | |
| 1252 | Contig 69 | | |
| 1253 | Contig 70 | | |
| 1254 | Contig 71 | | |
| 1255 | Contig 72 | | |
| 1256 | Contig 73 | | |
| 1257 | Contig 74 | | |
| 1258 | Contig 75 | | |
| 1259 | Contig 77 | | |
| 1260 | Contig 78 | | |
| 1261 | Contig 79 | | |
| 1262 | Contig 80 | | |
| 1263 | Contig 81 | | |
| 1264 | Contig 83 | | |
| 1265 | Contig 84 | | |
| 1266 | Contig 86 | | |
| 1267 | Contig 87 | | |
| 1268 | Contig 88 | | |
| 1269 | Contig 89 | | |
| 1270 | Contig 90 | | |
| 1271 | Contig 91 | | |
| 1272 | Contig 92 | | |
| 1273 | Contig 94 | | |
| 1274 | Contig 95 | | |
| 1275 | Contig 96 | | |
| 1276 | Contig 97 | | |
| 1277 | Contig 98 | | |

Table 3:
Clone names for NSCLC-SCLI and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1280 | Contig 38 | 1320 | Contig 82 |
| 1281 | Contig 39 | | |
| 1282 | Contig 40 | | |
| 1283 | Contig 41 | | |
| 1284 | Contig 42 | | |
| 1285 | Contig 43 | | |
| 1286 | Contig 44 | | |
| 1287 | Contig 45 | | |
| 1288 | Contig 46 | | |
| 1289 | Contig 47 | | |
| 1290 | Contig 48 | | |
| 1291 | Contig 49 | | |
| 1292 | Contig 51 | | |
| 1293 | Contig 52 | | |
| 1294 | Contig 53 | | |
| 1295 | Contig 54 | | |
| 1296 | Contig 55 | | |
| 1297 | Contig 56 | | |
| 1298 | Contig 57 | | |
| 1299 | Contig 58 | | |
| 1300 | Contig 59 | | |
| 1301 | Contig 60 | | |
| 1302 | Contig 62 | | |
| 1303 | Contig 63 | | |
| 1304 | Contig 64 | | |
| 1305 | Contig 65 | | |
| 1306 | Contig 66 | | |
| 1307 | Contig 67 | | |
| 1308 | Contig 68 | | |
| 1309 | Contig 69 | | |
| 1310 | Contig 70 | | |
| 1311 | Contig 72 | | |
| 1312 | Contig 73 | | |
| 1313 | Contig 75 | | |
| 1314 | Contig 76 | | |
| 1315 | Contig 77 | | |
| 1316 | Contig 78 | | |
| 1317 | Contig 79 | | |

Table 4A:
Clone names for NSCLC-SCL3-SCL4 and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1321 | Contig 94 | 1363 | Contig 136 |
| 1322 | Contig 95 | 1364 | Contig 137 |
| 1323 | Contig 96 | 1365 | Contig 138 |
| 1324 | Contig 97 | 1366 | Contig 139 |
| 1325 | Contig 98 | 1367 | Contig 140 |
| 1326 | Contig 99 | 1368 | Contig 141 |
| 1327 | Contig 100 | 1369 | Contig 142 |
| 1328 | Contig 101 | 1370 | Contig 143 |
| 1329 | Contig 102 | 1371 | Contig 144 |
| 1330 | Contig 103 | 1372 | Contig 145 |
| 1331 | Contig 104 | 1373 | Contig 146 |
| 1332 | Contig 105 | 1374 | Contig 147 |
| 1333 | Contig 106 | 1375 | Contig 148 |
| 1334 | Contig 107 | 1376 | Contig 149 |
| 1335 | Contig 108 | 1377 | Contig 150 |
| 1336 | Contig 109 | 1378 | Contig 151 |
| 1337 | Contig 110 | 1379 | Contig 152 |
| 1338 | Contig 111 | 1380 | Contig 153 |
| 1339 | Contig 112 | 1381 | Contig 154 |
| 1340 | Contig 113 | 1382 | Contig 155 |
| 1341 | Contig 114 | 1383 | Contig 156 |
| 1342 | Contig 115 | 1384 | Contig 157 |
| 1343 | Contig 116 | 1385 | Contig 158 |
| 1344 | Contig 117 | 1386 | Contig 159 |
| 1345 | Contig 118 | 1387 | Contig 160 |
| 1346 | Contig 119 | 1388 | Contig 161 |
| 1347 | Contig 120 | 1389 | Contig 162 |
| 1348 | Contig 121 | 1390 | Contig 163 |
| 1349 | Contig 122 | 1391 | Contig 164 |
| 1350 | Contig 123 | 1392 | Contig 165 |
| 1351 | Contig 124 | 1393 | Contig 166 |
| 1352 | Contig 125 | 1394 | Contig 167 |
| 1353 | Contig 126 | 1395 | Contig 168 |
| 1354 | Contig 127 | 1396 | Contig 169 |
| 1355 | Contig 128 | 1397 | Contig 170 |
| 1356 | Contig 129 | 1398 | Contig 171 |
| 1357 | Contig 130 | 1399 | Contig 172 |
| 1358 | Contig 131 | 1400 | Contig 173 |
| 1359 | Contig 132 | 1401 | Contig 174 |

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1360 | Contig 133 | 1402 | Contig 175 |
| 1361 | Contig 134 | 1403 | Contig 176 |
| 1362 | Contig 135 | | |

Table 4B:
Clone names for NSCLC-SCL3-SCL4 and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1404 | Contig 177 | | |
| 1405 | Contig 178 | | |
| 1406 | Contig 179 | | |
| 1407 | Contig 180 | | |
| 1408 | Contig 181 | | |
| 1409 | Contig 182 | | |
| 1410 | Contig 183 | | |
| 1411 | Contig 184 | | |
| 1412 | Contig 185 | | |
| 1413 | Contig 186 | | |
| 1414 | Contig 187 | | |

Table 5:
Clone names for SCLC-SQL1 and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1415 | Contig 17 | | |
| 1416 | Contig 18 | | |
| 1417 | Contig 20 | | |
| 1418 | Contig 23 | | |
| 1419 | Contig 24 | | |
| 1420 | Contig 25 | | |
| 1421 | Contig 26 | | |
| 1422 | Contig 27 | | |
| 1423 | Contig 28 | | |
| 1424 | Contig 29 | | |
| 1425 | Contig 30 | | |
| 1426 | Contig 31 | | |
| 1427 | Contig 20 | | |
| 1428 | Contig 21 | | |
| 1429 | Contig 22 | | |
| 1430 | Contig 23 | | |
| 1431 | Contig 24 | | |
| 1432 | Contig 25 | | |
| 1433 | Contig 26 | | |
| 1434 | Contig 27 | | |
| 1435 | Contig 28 | | |
| 1436 | Contig 29 | | |
| 1437 | Contig 30 | | |
| 1438 | Contig 31 | | |
| 1439 | Contig 32 | | |
| 1440 | Contig 33 | | |
| 1441 | Contig 34 | | |
| 1442 | Contig 35 | | |
| 1443 | Contig 36 | | |
| 1444 | Contig 37 | | |
| 1445 | Contig 38 | | |

Table 6A:
Clone names for SCLC-SCL3-SCL4 and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1446 | Contig 116 | 1488 | Contig 160 |
| 1447 | Contig 117 | 1489 | Contig 161 |
| 1448 | Contig 118 | 1490 | Contig 162 |
| 1449 | Contig 119 | 1491 | Contig 163 |
| 1450 | Contig 120 | 1492 | Contig 164 |
| 1451 | Contig 122 | 1493 | Contig 165 |
| 1452 | Contig 123 | 1494 | Contig 166 |
| 1453 | Contig 124 | 1495 | Contig 167 |
| 1454 | Contig 125 | 1496 | Contig 168 |
| 1455 | Contig 126 | 1497 | Contig 169 |
| 1456 | Contig 127 | 1498 | Contig 170 |
| 1457 | Contig 128 | 1499 | Contig 171 |
| 1458 | Contig 129 | 1500 | Contig 172 |
| 1459 | Contig 130 | 1501 | Contig 173 |
| 1460 | Contig 131 | 1502 | Contig 174 |
| 1461 | Contig 132 | 1503 | Contig 175 |
| 1462 | Contig 133 | 1504 | Contig 176 |
| 1463 | Contig 135 | 1505 | Contig 177 |
| 1464 | Contig 136 | 1506 | Contig 178 |
| 1465 | Contig 137 | 1507 | Contig 179 |
| 1466 | Contig 138 | 1508 | Contig 181 |
| 1467 | Contig 139 (L985P) | 1509 | Contig 182 |
| 1468 | Contig 140 | 1510 | Contig 183 |
| 1469 | Contig 141 | 1511 | Contig 184 |
| 1470 | Contig 142 | 1512 | Contig 185 |
| 1471 | Contig 143 | 1513 | Contig 186 |
| 1472 | Contig 144 | 1514 | Contig 187 |
| 1473 | Contig 145 | 1515 | Contig 189 |
| 1474 | Contig 146 | 1516 | Contig 190 |
| 1475 | Contig 147 | 1517 | Contig 191 |
| 1476 | Contig 148 | 1518 | Contig 192 |
| 1477 | Contig 149 | 1519 | Contig 193 |
| 1478 | Contig 150 | 1520 | Contig 194 |
| 1479 | Contig 151 | 1521 | Contig 195 |
| 1480 | Contig 152 | 1522 | Contig 196 |
| 1481 | Contig 153 | 1523 | Contig 197 |
| 1482 | Contig 154 | 1524 | Contig 198 |
| 1483 | Contig 155 | 1525 | Contig 199 |
| 1484 | Contig 156 | 1526 | Contig 200 |

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1485 | Contig 157 | 1527 | Contig 201 |
| 1486 | Contig 158 | 1528 | Contig 202 |
| 1487 | Contig 159 | | |

20250121 15:30:00

Table 6B:
Clone names for SCLC-SCL3-SCL4 and corresponding SEQ ID NOs

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|------------------|-------------------------|------------------|-------------------------|
| 1529 | Contig 203 | | |
| 1530 | Contig 204 | | |
| 1531 | Contig 205 | | |
| 1532 | Contig 206 | | |
| 1533 | Contig 207 | | |
| 1534 | Contig 208 | | |
| 1535 | Contig 209 | | |
| 1536 | Contig 210 | | |
| 1537 | Contig 211 | | |
| 1538 | Contig 212 | | |
| 1539 | Contig 213 | | |
| 1540 | Contig 214 | | |
| 1541 | Contig 215 | | |
| 1542 | Contig 216 | | |
| 1543 | Contig 217 | | |
| 1544 | Contig 218 | | |
| 1545 | Contig 219 | | |
| 1546 | Contig 220 | | |
| 1547 | Contig 221 | | |
| 1548 | Contig 222 | | |
| 1549 | Contig 223 | | |
| 1550 | Contig 224 | | |
| 1551 | Contig 225 | | |
| 1552 | Contig 226 | | |
| 1553 | Contig 227 | | |
| 1554 | Contig 228 | | |
| 1555 | Contig 229 | | |
| 1556 | Contig 230 | | |
| 1557 | Contig 231 | | |
| 1558 | Contig 232 | | |
| 1559 | Contig 233 | | |
| 1560 | Contig 234 | | |
| 1561 | Contig 235 | | |
| 1562 | Contig 236 | | |
| 1563 | Contig 237 | | |

Table 7.

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1564 | R0124:E05 | 1609 | R0129:D09 |
| 1565 | R0124:E06 | 1610 | R0129:D10 |
| 1566 | R0124:E08 | 1611 | R0129:D11 |
| 1567 | R0124:F07 | 1612 | R0129:E02 |
| 1568 | R0124:F08 | 1613 | R0129:E03 |
| 1569 | R0124:F09 | 1614 | R0129:E04 |
| 1570 | R0124:G04 | 1615 | R0129:E05 |
| 1571 | R0129:A02 | 1616 | R0129:E06 |
| 1572 | R0129:A03 | 1617 | R0129:E07 |
| 1573 | R0129:A06 | 1618 | R0129:E08 |
| 1574 | R0129:A07 | 1619 | R0129:E09 |
| 1575 | R0129:A08 | 1620 | R0129:E11 |
| 1576 | R0129:A09 | 1621 | R0129:E12 |
| 1577 | R0129:A10 | 1622 | R0129:F01 |
| 1578 | R0129:A11 | 1623 | R0129:F02 |
| 1579 | R0129:A12 | 1624 | R0129:F03 |
| 1580 | R0129:B02 | 1625 | R0129:F04 |
| 1581 | R0129:B03 | 1626 | R0129:F06 |
| 1582 | R0129:B04 | 1627 | R0129:F07 |
| 1583 | R0129:B05 | 1628 | R0129:F08 |
| 1584 | R0129:B06 | 1629 | R0129:F09 |
| 1585 | R0129:B07 | 1630 | R0129:F10 |
| 1586 | R0129:B08 | 1631 | R0129:F11 |
| 1587 | R0129:B09 | 1632 | R0129:F12 |
| 1588 | R0129:B10 | 1633 | R0129:G01 |
| 1589 | R0129:B11 | 1634 | R0129:G02 |
| 1590 | R0129:B12 | 1635 | R0129:G03 |
| 1591 | R0129:C01 | 1636 | R0129:G04 |
| 1592 | R0129:C02 | 1637 | R0129:G05 |
| 1593 | R0129:C03 | 1638 | R0129:G06 |
| 1594 | R0129:C04 | 1639 | R0129:G07 |
| 1595 | R0129:C06 | 1640 | R0129:G08 |
| 1596 | R0129:C07 | 1641 | R0129:G09 |
| 1597 | R0129:C08 | 1642 | R0129:G10 |
| 1598 | R0129:C09 | 1643 | R0129:G11 |
| 1599 | R0129:C10 | 1644 | R0129:G12 |
| 1600 | R0129:C11 | 1645 | R0129:H01 |
| 1601 | R0129:C12 | 1646 | R0129:H02 |
| 1602 | R0129:D01 | 1647 | R0129:H03 |
| 1603 | R0129:D03 | 1648 | R0129:H04 |

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1604 | R0129:D04 | 1649 | R0129:H05 |
| 1605 | R0129:D05 | 1650 | R0129:H08 |
| 1606 | R0129:D06 | 1651 | R0129:H09 |
| 1607 | R0129:D07 | 1652 | R0129:H10 |
| 1608 | R0129:D08 | 1653 | R0129:H11 |

20250304 16:43:43

Table 8.

| SEQ ID NO | CLONE IDENTIFIER | SEQ ID NO | CLONE IDENTIFIER |
|-----------|------------------|-----------|------------------|
| 1654 | 26484 | | |
| 1655 | 26496 | | |
| 1656 | 26517 | | |
| 1657 | 26531 | | |
| 1658 | 26022 | | |
| 1659 | 26026 | | |
| 1660 | 26810 | | |
| 1661 | 26815 | | |
| 1662 | 26869 | | |
| 1663 | 26883 | | |
| 1664 | 26902 | | |

SEQ ID NO:1667 is the protein sequence of expressed recombinant

5 L7548S.

SEQ ID NO:1668 is the cDNA sequence of expressed recombinant L7548S.

SEQ ID NO:1669 is the extended cDNA sequence of clone #18971

(L801P).

10 SEQ ID NO:1670 is the amino acid sequence of open reading frame ORF4 encoded by SEQ ID NO:1669.

SEQ ID NO:1671 is the amino acid sequence of open reading frame ORF5 encoded by SEQ ID NO:1669.

SEQ ID NO:1672 is the amino acid sequence of open reading frame ORF6 encoded by SEQ ID NO:1669.

15 SEQ ID NO:1673 is the amino acid sequence of open reading frame ORF7 encoded by SEQ ID NO:1669.

SEQ ID NO:1674 is the amino acid sequence of open reading frame ORF8 encoded by SEQ ID NO:1669.

20 SEQ ID NO:1675 is the amino acid sequence of open reading frame ORF9 encoded by SEQ ID NO:1669.

SEQ ID NO:1676 is the extended cDNA for contig 139 (SEQ ID NO:1467), also known as L985P.

SEQ ID NO:1677 is the L985P amino acid sequence encoded by SEQ ID NO: 1676.

SEQ ID NO: 1678 is the amino acid sequence of open reading frame ORF5X of SEQ ID NO:1669.

- 5 SEQ ID NO: 1679 is the amino acid sequence of an open reading frame for contig 139 (SEQ ID NO:1467).

DETAILED DESCRIPTION OF THE INVENTION

As noted above, the present invention is generally directed to compositions and methods for using the compositions, for example in the therapy and diagnosis of cancer, such as lung cancer. Certain illustrative compositions described herein include lung tumor polypeptides, polynucleotides encoding such polypeptides, binding agents such as antibodies, antigen presenting cells (APCs) and/or immune system cells (*e.g.*, T cells). A "lung tumor protein," as the term is used herein, refers generally to a protein that is expressed in lung tumor cells at a level that is at least two fold, and preferably at least five fold, greater than the level of expression in a normal tissue, as determined using a representative assay provided herein. Certain lung tumor proteins are tumor proteins that react detectably (within an immunoassay, such as an ELISA or Western blot) with antisera of a patient afflicted with lung cancer.

20 Therefore, in accordance with the above, and as described further below, the present invention provides illustrative polynucleotide compositions having sequences set forth in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319,

1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 illustrative polypeptide compositions having amino acid sequences set forth in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809, 827, 1670-1675 and 1677-1679, antibody compositions capable of binding such polypeptides, and numerous additional embodiments employing such compositions, for example in the detection, diagnosis and/or therapy of human lung cancer.

10 POLYNUCLEOTIDE COMPOSITIONS

As used herein, the terms "DNA segment" and "polynucleotide" refer to a DNA molecule that has been isolated free of total genomic DNA of a particular species. Therefore, a DNA segment encoding a polypeptide refers to a DNA segment that contains one or more coding sequences yet is substantially isolated away from, or purified free from, total genomic DNA of the species from which the DNA segment is obtained. Included within the terms "DNA segment" and "polynucleotide" are DNA segments and smaller fragments of such segments, and also recombinant vectors, including, for example, plasmids, cosmids, phagemids, phage, viruses, and the like.

As will be understood by those skilled in the art, the DNA segments of this invention can include genomic sequences, extra-genomic and plasmid-encoded sequences and smaller engineered gene segments that express, or may be adapted to express, proteins, polypeptides, peptides and the like. Such segments may be naturally isolated, or modified synthetically by the hand of man.

"Isolated," as used herein, means that a polynucleotide is substantially away from other coding sequences, and that the DNA segment does not contain large portions of unrelated coding DNA, such as large chromosomal fragments or other functional genes or polypeptide coding regions. Of course, this refers to the DNA segment as originally

isolated, and does not exclude genes or coding regions later added to the segment by the hand of man.

As will be recognized by the skilled artisan, polynucleotides may be single-stranded (coding or antisense) or double-stranded, and may be DNA (genomic, cDNA or synthetic) or RNA molecules. RNA molecules include HnRNA molecules, which contain introns and correspond to a DNA molecule in a one-to-one manner, and mRNA molecules, which do not contain introns. Additional coding or non-coding sequences may, but need not, be present within a polynucleotide of the present invention, and a polynucleotide may, but need not, be linked to other molecules and/or support materials.

Polynucleotides may comprise a native sequence (*i.e.*, an endogenous sequence that encodes a lung tumor protein or a portion thereof) or may comprise a variant, or a biological or antigenic functional equivalent of such a sequence. Polynucleotide variants may contain one or more substitutions, additions, deletions and/or insertions, as further described below, preferably such that the immunogenicity of the encoded polypeptide is not diminished, relative to a native tumor protein. The effect on the immunogenicity of the encoded polypeptide may generally be assessed as described herein. The term “variants” also encompasses homologous genes of xenogenic origin.

When comparing polynucleotide or polypeptide sequences, two sequences are said to be “identical” if the sequence of nucleotides or amino acids in the two sequences is the same when aligned for maximum correspondence, as described below. Comparisons between two sequences are typically performed by comparing the sequences over a comparison window to identify and compare local regions of sequence similarity. A “comparison window” as used herein, refers to a segment of at least about 20 contiguous positions, usually 30 to about 75, 40 to about 50, in which a sequence may be compared to a reference sequence of the same number of contiguous positions after the two sequences are optimally aligned.

Optimal alignment of sequences for comparison may be conducted using the Megalign program in the Lasergene suite of bioinformatics software (DNASTAR, Inc., Madison, WI), using default parameters. This program embodies several alignment

- schemes described in the following references: Dayhoff, M.O. (1978) A model of evolutionary change in proteins – Matrices for detecting distant relationships. In Dayhoff, M.O. (ed.) *Atlas of Protein Sequence and Structure*, National Biomedical Research Foundation, Washington DC Vol. 5, Suppl. 3, pp. 345-358; Hein J. (1990) Unified
- 5 Approach to Alignment and Phylogenesis pp. 626-645 *Methods in Enzymology* vol. 183, Academic Press, Inc., San Diego, CA; Higgins, D.G. and Sharp, P.M. (1989) *CABIOS* 5:151-153; Myers, E.W. and Muller W. (1988) *CABIOS* 4:11-17; Robinson, E.D. (1971) *Comb. Theor* 11:105; Santou, N. Nes, M. (1987) *Mol. Biol. Evol.* 4:406-425; Sneath, P.H.A. and Sokal, R.R. (1973) *Numerical Taxonomy – the Principles and Practice of*
- 10 *Numerical Taxonomy*, Freeman Press, San Francisco, CA; Wilbur, W.J. and Lipman, D.J. (1983) *Proc. Natl. Acad., Sci. USA* 80:726-730.

Alternatively, optimal alignment of sequences for comparison may be conducted by the local identity algorithm of Smith and Waterman (1981) *Add. APL. Math* 2:482, by the identity alignment algorithm of Needleman and Wunsch (1970) *J. Mol. Biol.*

15 48:443, by the search for similarity methods of Pearson and Lipman (1988) *Proc. Natl. Acad. Sci. USA* 85: 2444, by computerized implementations of these algorithms (GAP, BESTFIT, BLAST, FASTA, and TFASTA in the Wisconsin Genetics Software Package, Genetics Computer Group (GCG), 575 Science Dr., Madison, WI), or by inspection.

One preferred example of algorithms that are suitable for determining

20 percent sequence identity and sequence similarity are the BLAST and BLAST 2.0 algorithms, which are described in Altschul *et al.* (1977) *Nucl. Acids Res.* 25:3389-3402 and Altschul *et al.* (1990) *J. Mol. Biol.* 215:403-410, respectively. BLAST and BLAST 2.0 can be used, for example with the parameters described herein, to determine percent sequence identity for the polynucleotides and polypeptides of the invention. Software for

25 performing BLAST analyses is publicly available through the National Center for Biotechnology Information. In one illustrative example, cumulative scores can be calculated using, for nucleotide sequences, the parameters M (reward score for a pair of matching residues; always >0) and N (penalty score for mismatching residues; always <0). For amino acid sequences, a scoring matrix can be used to calculate the cumulative score.

Extension of the word hits in each direction are halted when: the cumulative alignment score falls off by the quantity X from its maximum achieved value; the cumulative score goes to zero or below, due to the accumulation of one or more negative-scoring residue alignments; or the end of either sequence is reached. The BLAST algorithm parameters W, T and X determine the sensitivity and speed of the alignment. The BLASTN program (for nucleotide sequences) uses as defaults a wordlength (W) of 11, and expectation (E) of 10, and the BLOSUM62 scoring matrix (see Henikoff and Henikoff (1989) *Proc. Natl. Acad. Sci. USA* 89:10915) alignments, (B) of 50, expectation (E) of 10, M=5, N=-4 and a comparison of both strands.

Preferably, the "percentage of sequence identity" is determined by comparing two optimally aligned sequences over a window of comparison of at least 20 positions, wherein the portion of the polynucleotide or polypeptide sequence in the comparison window may comprise additions or deletions (*i.e.*, gaps) of 20 percent or less, usually 5 to 15 percent, or 10 to 12 percent, as compared to the reference sequences (which does not comprise additions or deletions) for optimal alignment of the two sequences. The percentage is calculated by determining the number of positions at which the identical nucleic acid bases or amino acid residue occurs in both sequences to yield the number of matched positions, dividing the number of matched positions by the total number of positions in the reference sequence (*i.e.*, the window size) and multiplying the results by 100 to yield the percentage of sequence identity.

Therefore, the present invention encompasses polynucleotide and polypeptide sequences having substantial identity to the sequences disclosed herein, for example those comprising at least 50% sequence identity, preferably at least 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% or higher, sequence identity compared to a polynucleotide or polypeptide sequence of this invention using the methods described herein, (*e.g.*, BLAST analysis using standard parameters, as described below). One skilled in this art will recognize that these values can be appropriately adjusted to determine corresponding identity of proteins encoded by two nucleotide

sequences by taking into account codon degeneracy, amino acid similarity, reading frame positioning and the like.

In additional embodiments, the present invention provides isolated polynucleotides and polypeptides comprising various lengths of contiguous stretches of
 5 sequence identical to or complementary to one or more of the sequences disclosed herein. For example, polynucleotides are provided by this invention that comprise at least about 15, 20, 30, 40, 50, 75, 100, 150, 200, 300, 400, 500 or 1000 or more contiguous nucleotides of one or more of the sequences disclosed herein as well as all intermediate lengths there between. It will be readily understood that "intermediate lengths", in this context, means
 10 any length between the quoted values, such as 16, 17, 18, 19, *etc.*; 21, 22, 23, *etc.*; 30, 31, 32, *etc.*; 50, 51, 52, 53, *etc.*; 100, 101, 102, 103, *etc.*; 150, 151, 152, 153, *etc.*; including all integers through 200-500; 500-1,000, and the like.

The polynucleotides of the present invention, or fragments thereof, regardless of the length of the coding sequence itself, may be combined with other DNA
 15 sequences, such as promoters, polyadenylation signals, additional restriction enzyme sites, multiple cloning sites, other coding segments, and the like, such that their overall length may vary considerably. It is therefore contemplated that a nucleic acid fragment of almost any length may be employed, with the total length preferably being limited by the ease of preparation and use in the intended recombinant DNA protocol. For example, illustrative
 20 DNA segments with total lengths of about 10,000, about 5000, about 3000, about 2,000, about 1,000, about 500, about 200, about 100, about 50 base pairs in length, and the like, (including all intermediate lengths) are contemplated to be useful in many implementations of this invention.

In other embodiments, the present invention is directed to polynucleotides
 25 that are capable of hybridizing under moderately stringent conditions to a polynucleotide sequence provided herein, or a fragment thereof, or a complementary sequence thereof. Hybridization techniques are well known in the art of molecular biology. For purposes of illustration, suitable moderately stringent conditions for testing the hybridization of a polynucleotide of this invention with other polynucleotides include prewashing in a

solution of 5 X SSC, 0.5% SDS, 1.0 mM EDTA (pH 8.0); hybridizing at 50°C-65°C, 5 X SSC, overnight; followed by washing twice at 65°C for 20 minutes with each of 2X, 0.5X and 0.2X SSC containing 0.1% SDS.

Moreover, it will be appreciated by those of ordinary skill in the art that, as a result of the degeneracy of the genetic code, there are many nucleotide sequences that encode a polypeptide as described herein. Some of these polynucleotides bear minimal homology to the nucleotide sequence of any native gene. Nonetheless, polynucleotides that vary due to differences in codon usage are specifically contemplated by the present invention. Further, alleles of the genes comprising the polynucleotide sequences provided herein are within the scope of the present invention. Alleles are endogenous genes that are altered as a result of one or more mutations, such as deletions, additions and/or substitutions of nucleotides. The resulting mRNA and protein may, but need not, have an altered structure or function. Alleles may be identified using standard techniques (such as hybridization, amplification and/or database sequence comparison).

15 **PROBES AND PRIMERS**

In other embodiments of the present invention, the polynucleotide sequences provided herein can be advantageously used as probes or primers for nucleic acid hybridization. As such, it is contemplated that nucleic acid segments that comprise a sequence region of at least about 15 nucleotide long contiguous sequence that has the same sequence as, or is complementary to, a 15 nucleotide long contiguous sequence disclosed herein will find particular utility. Longer contiguous identical or complementary sequences, *e.g.*, those of about 20, 30, 40, 50, 100, 200, 500, 1000 (including all intermediate lengths) and even up to full length sequences will also be of use in certain embodiments.

25 The ability of such nucleic acid probes to specifically hybridize to a sequence of interest will enable them to be of use in detecting the presence of complementary sequences in a given sample. However, other uses are also envisioned,

such as the use of the sequence information for the preparation of mutant species primers, or primers for use in preparing other genetic constructions.

Polynucleotide molecules having sequence regions consisting of contiguous nucleotide stretches of 10-14, 15-20, 30, 50, or even of 100-200 nucleotides or so (including intermediate lengths as well), identical or complementary to a polynucleotide sequence disclosed herein, are particularly contemplated as hybridization probes for use in, *e.g.*, Southern and Northern blotting. This would allow a gene product, or fragment thereof, to be analyzed, both in diverse cell types and also in various bacterial cells. The total size of fragment, as well as the size of the complementary stretch(es), will ultimately depend on the intended use or application of the particular nucleic acid segment. Smaller fragments will generally find use in hybridization embodiments, wherein the length of the contiguous complementary region may be varied, such as between about 15 and about 100 nucleotides, but larger contiguous complementarity stretches may be used, according to the length complementary sequences one wishes to detect.

The use of a hybridization probe of about 15-25 nucleotides in length allows the formation of a duplex molecule that is both stable and selective. Molecules having contiguous complementary sequences over stretches greater than 15 bases in length are generally preferred, though, in order to increase stability and selectivity of the hybrid, and thereby improve the quality and degree of specific hybrid molecules obtained. One will generally prefer to design nucleic acid molecules having gene-complementary stretches of 15 to 25 contiguous nucleotides, or even longer where desired.

Hybridization probes may be selected from any portion of any of the sequences disclosed herein. All that is required is to review the sequence set forth in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669

and 1676, or to any continuous portion of the sequence, from about 15-25 nucleotides in length up to and including the full length sequence, that one wishes to utilize as a probe or primer. The choice of probe and primer sequences may be governed by various factors. For example, one may wish to employ primers from towards the termini of the total
5 sequence.

Small polynucleotide segments or fragments may be readily prepared by, for example, directly synthesizing the fragment by chemical means, as is commonly practiced using an automated oligonucleotide synthesizer. Also, fragments may be obtained by application of nucleic acid reproduction technology, such as the PCRTM technology of U. S.
10 Patent 4,683,202 (incorporated herein by reference), by introducing selected sequences into recombinant vectors for recombinant production, and by other recombinant DNA techniques generally known to those of skill in the art of molecular biology.

The nucleotide sequences of the invention may be used for their ability to selectively form duplex molecules with complementary stretches of the entire gene or gene
15 fragments of interest. Depending on the application envisioned, one will typically desire to employ varying conditions of hybridization to achieve varying degrees of selectivity of probe towards target sequence. For applications requiring high selectivity, one will typically desire to employ relatively stringent conditions to form the hybrids, *e.g.*, one will select relatively low salt and/or high temperature conditions, such as provided by a salt
20 concentration of from about 0.02 M to about 0.15 M salt at temperatures of from about 50°C to about 70°C. Such selective conditions tolerate little, if any, mismatch between the probe and the template or target strand, and would be particularly suitable for isolating related sequences.

Of course, for some applications, for example, where one desires to prepare
25 mutants employing a mutant primer strand hybridized to an underlying template, less stringent (reduced stringency) hybridization conditions will typically be needed in order to allow formation of the heteroduplex. In these circumstances, one may desire to employ salt conditions such as those of from about 0.15 M to about 0.9 M salt, at temperatures ranging from about 20°C to about 55°C. Cross-hybridizing species can thereby be readily

identified as positively hybridizing signals with respect to control hybridizations. In any case, it is generally appreciated that conditions can be rendered more stringent by the addition of increasing amounts of formamide, which serves to destabilize the hybrid duplex in the same manner as increased temperature. Thus, hybridization conditions can be readily manipulated, and thus will generally be a method of choice depending on the desired results.

POLYNUCLEOTIDE IDENTIFICATION AND CHARACTERIZATION

Polynucleotides may be identified, prepared and/or manipulated using any of a variety of well established techniques. For example, a polynucleotide may be identified, as described in more detail below, by screening a microarray of cDNAs for tumor-associated expression (*i.e.*, expression that is at least two fold greater in a tumor than in normal tissue, as determined using a representative assay provided herein). Such screens may be performed, for example, using a Synteni microarray (Palo Alto, CA) according to the manufacturer's instructions (and essentially as described by Schena *et al.*, *Proc. Natl. Acad. Sci. USA* 93:10614-10619, 1996 and Heller *et al.*, *Proc. Natl. Acad. Sci. USA* 94:2150-2155, 1997). Alternatively, polynucleotides may be amplified from cDNA prepared from cells expressing the proteins described herein, such as lung tumor cells. Such polynucleotides may be amplified via polymerase chain reaction (PCR). For this approach, sequence-specific primers may be designed based on the sequences provided herein, and may be purchased or synthesized.

An amplified portion of a polynucleotide of the present invention may be used to isolate a full length gene from a suitable library (*e.g.*, a lung tumor cDNA library) using well known techniques. Within such techniques, a library (cDNA or genomic) is screened using one or more polynucleotide probes or primers suitable for amplification. Preferably, a library is size-selected to include larger molecules. Random primed libraries may also be preferred for identifying 5' and upstream regions of genes. Genomic libraries are preferred for obtaining introns and extending 5' sequences.

For hybridization techniques, a partial sequence may be labeled (*e.g.*, by nick-translation or end-labeling with ^{32}P) using well known techniques. A bacterial or bacteriophage library is then generally screened by hybridizing filters containing denatured bacterial colonies (or lawns containing phage plaques) with the labeled probe (*see* 5 Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY, 1989). Hybridizing colonies or plaques are selected and expanded, and the DNA is isolated for further analysis. cDNA clones may be analyzed to determine the amount of additional sequence by, for example, PCR using a primer from the partial sequence and a primer from the vector. Restriction maps and 10 partial sequences may be generated to identify one or more overlapping clones. The complete sequence may then be determined using standard techniques, which may involve generating a series of deletion clones. The resulting overlapping sequences can then be assembled into a single contiguous sequence. A full length cDNA molecule can be generated by ligating suitable fragments, using well known techniques.

15 Alternatively, there are numerous amplification techniques for obtaining a full length coding sequence from a partial cDNA sequence. Within such techniques, amplification is generally performed via PCR. Any of a variety of commercially available kits may be used to perform the amplification step. Primers may be designed using, for example, software well known in the art. Primers are preferably 22-30 nucleotides in 20 length, have a GC content of at least 50% and anneal to the target sequence at temperatures of about 68°C to 72°C. The amplified region may be sequenced as described above, and overlapping sequences assembled into a contiguous sequence.

One such amplification technique is inverse PCR (*see* Triglia *et al.*, *Nucl. Acids Res.* 16:8186, 1988), which uses restriction enzymes to generate a fragment in the 25 known region of the gene. The fragment is then circularized by intramolecular ligation and used as a template for PCR with divergent primers derived from the known region. Within an alternative approach, sequences adjacent to a partial sequence may be retrieved by amplification with a primer to a linker sequence and a primer specific to a known region. The amplified sequences are typically subjected to a second round of amplification with the

same linker primer and a second primer specific to the known region. A variation on this procedure, which employs two primers that initiate extension in opposite directions from the known sequence, is described in WO 96/38591. Another such technique is known as "rapid amplification of cDNA ends" or RACE. This technique involves the use of an
 5 internal primer and an external primer, which hybridizes to a polyA region or vector sequence, to identify sequences that are 5' and 3' of a known sequence. Additional techniques include capture PCR (Lagerstrom *et al.*, *PCR Methods Applic.* 1:111-19, 1991) and walking PCR (Parker *et al.*, *Nucl. Acids. Res.* 19:3055-60, 1991). Other methods employing amplification may also be employed to obtain a full length cDNA sequence.

10 In certain instances, it is possible to obtain a full length cDNA sequence by analysis of sequences provided in an expressed sequence tag (EST) database, such as that available from GenBank. Searches for overlapping ESTs may generally be performed using well known programs (*e.g.*, NCBI BLAST searches), and such ESTs may be used to generate a contiguous full length sequence. Full length DNA sequences may also be
 15 obtained by analysis of genomic fragments.

POLYNUCLEOTIDE EXPRESSION IN HOST CELLS

In other embodiments of the invention, polynucleotide sequences or fragments thereof which encode polypeptides of the invention, or fusion proteins or functional equivalents thereof, may be used in recombinant DNA molecules to direct
 20 expression of a polypeptide in appropriate host cells. Due to the inherent degeneracy of the genetic code, other DNA sequences that encode substantially the same or a functionally equivalent amino acid sequence may be produced and these sequences may be used to clone and express a given polypeptide.

As will be understood by those of skill in the art, it may be advantageous in
 25 some instances to produce polypeptide-encoding nucleotide sequences possessing non-naturally occurring codons. For example, codons preferred by a particular prokaryotic or eukaryotic host can be selected to increase the rate of protein expression or to produce a

recombinant RNA transcript having desirable properties, such as a half-life which is longer than that of a transcript generated from the naturally occurring sequence.

Moreover, the polynucleotide sequences of the present invention can be engineered using methods generally known in the art in order to alter polypeptide encoding sequences for a variety of reasons, including but not limited to, alterations which modify the cloning, processing, and/or expression of the gene product. For example, DNA shuffling by random fragmentation and PCR reassembly of gene fragments and synthetic oligonucleotides may be used to engineer the nucleotide sequences. In addition, site-directed mutagenesis may be used to insert new restriction sites, alter glycosylation patterns, change codon preference, produce splice variants, or introduce mutations, and so forth.

In another embodiment of the invention, natural, modified, or recombinant nucleic acid sequences may be ligated to a heterologous sequence to encode a fusion protein. For example, to screen peptide libraries for inhibitors of polypeptide activity, it may be useful to encode a chimeric protein that can be recognized by a commercially available antibody. A fusion protein may also be engineered to contain a cleavage site located between the polypeptide-encoding sequence and the heterologous protein sequence, so that the polypeptide may be cleaved and purified away from the heterologous moiety.

Sequences encoding a desired polypeptide may be synthesized, in whole or in part, using chemical methods well known in the art (see Caruthers, M. H. *et al.* (1980) *Nucl. Acids Res. Symp. Ser.* 215-223, Horn, T. *et al.* (1980) *Nucl. Acids Res. Symp. Ser.* 225-232). Alternatively, the protein itself may be produced using chemical methods to synthesize the amino acid sequence of a polypeptide, or a portion thereof. For example, peptide synthesis can be performed using various solid-phase techniques (Roberge, J. Y. *et al.* (1995) *Science* 269:202-204) and automated synthesis may be achieved, for example, using the ABI 431A Peptide Synthesizer (Perkin Elmer, Palo Alto, CA).

A newly synthesized peptide may be substantially purified by preparative high performance liquid chromatography (*e.g.*, Creighton, T. (1983) *Proteins, Structures and Molecular Principles*, WH Freeman and Co., New York, N.Y.) or other comparable

techniques available in the art. The composition of the synthetic peptides may be confirmed by amino acid analysis or sequencing (*e.g.*, the Edman degradation procedure). Additionally, the amino acid sequence of a polypeptide, or any part thereof, may be altered during direct synthesis and/or combined using chemical methods with sequences from
 5 other proteins, or any part thereof, to produce a variant polypeptide.

In order to express a desired polypeptide, the nucleotide sequences encoding the polypeptide, or functional equivalents, may be inserted into appropriate expression vector, *i.e.*, a vector which contains the necessary elements for the transcription and translation of the inserted coding sequence. Methods which are well known to those
 10 skilled in the art may be used to construct expression vectors containing sequences encoding a polypeptide of interest and appropriate transcriptional and translational control elements. These methods include *in vitro* recombinant DNA techniques, synthetic techniques, and *in vivo* genetic recombination. Such techniques are described in Sambrook, J. *et al.* (1989) *Molecular Cloning, A Laboratory Manual*, Cold Spring Harbor Press, Plainview, N.Y., and Ausubel, F. M. *et al.* (1989) *Current Protocols in Molecular Biology*,
 15 John Wiley & Sons, New York, N.Y.

A variety of expression vector/host systems may be utilized to contain and express polynucleotide sequences. These include, but are not limited to, microorganisms such as bacteria transformed with recombinant bacteriophage, plasmid, or cosmid DNA
 20 expression vectors; yeast transformed with yeast expression vectors; insect cell systems infected with virus expression vectors (*e.g.*, baculovirus); plant cell systems transformed with virus expression vectors (*e.g.*, cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or with bacterial expression vectors (*e.g.*, Ti or pBR322 plasmids); or animal cell systems.

25 The "control elements" or "regulatory sequences" present in an expression vector are those non-translated regions of the vector--enhancers, promoters, 5' and 3' untranslated regions--which interact with host cellular proteins to carry out transcription and translation. Such elements may vary in their strength and specificity. Depending on the vector system and host utilized, any number of suitable transcription and translation

elements, including constitutive and inducible promoters, may be used. For example, when cloning in bacterial systems, inducible promoters such as the hybrid lacZ promoter of the PBLUESCRIPT phagemid (Stratagene, La Jolla, Calif.) or PSFORT1 plasmid (Gibco BRL, Gaithersburg, MD) and the like may be used. In mammalian cell systems, promoters from

5 mammalian genes or from mammalian viruses are generally preferred. If it is necessary to generate a cell line that contains multiple copies of the sequence encoding a polypeptide, vectors based on SV40 or EBV may be advantageously used with an appropriate selectable marker.

In bacterial systems, a number of expression vectors may be selected

10 depending upon the use intended for the expressed polypeptide. For example, when large quantities are needed, for example for the induction of antibodies, vectors which direct high level expression of fusion proteins that are readily purified may be used. Such vectors include, but are not limited to, the multifunctional *E. coli* cloning and expression vectors such as BLUESCRIPT (Stratagene), in which the sequence encoding the polypeptide of

15 interest may be ligated into the vector in frame with sequences for the amino-terminal Met and the subsequent 7 residues of β -galactosidase so that a hybrid protein is produced; pIN vectors (Van Heeke, G. and S. M. Schuster (1989) *J. Biol. Chem.* 264:5503-5509); and the like. pGEX Vectors (Promega, Madison, Wis.) may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such

20 fusion proteins are soluble and can easily be purified from lysed cells by adsorption to glutathione-agarose beads followed by elution in the presence of free glutathione. Proteins made in such systems may be designed to include heparin, thrombin, or factor XA protease cleavage sites so that the cloned polypeptide of interest can be released from the GST moiety at will.

25 In the yeast, *Saccharomyces cerevisiae*, a number of vectors containing constitutive or inducible promoters such as alpha factor, alcohol oxidase, and PGH may be used. For reviews, see Ausubel *et al.* (supra) and Grant *et al.* (1987) *Methods Enzymol.* 153:516-544.

In cases where plant expression vectors are used, the expression of sequences encoding polypeptides may be driven by any of a number of promoters. For example, viral promoters such as the 35S and 19S promoters of CaMV may be used alone or in combination with the omega leader sequence from TMV (Takamatsu, N. (1987) *EMBO J.* 6:307-311. Alternatively, plant promoters such as the small subunit of RUBISCO or heat shock promoters may be used (Coruzzi, G. *et al.* (1984) *EMBO J.* 3:1671-1680; Broglie, R. *et al.* (1984) *Science* 224:838-843; and Winter, J. *et al.* (1991) *Results Probl. Cell Differ.* 17:85-105). These constructs can be introduced into plant cells by direct DNA transformation or pathogen-mediated transfection. Such techniques are described in a number of generally available reviews (see, for example, Hobbs, S. or Murry, L. E. in McGraw Hill Yearbook of Science and Technology (1992) McGraw Hill, New York, N.Y.; pp. 191-196).

An insect system may also be used to express a polypeptide of interest. For example, in one such system, Autographa californica nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes in *Spodoptera frugiperda* cells or in *Trichoplusia* larvae. The sequences encoding the polypeptide may be cloned into a non-essential region of the virus, such as the polyhedrin gene, and placed under control of the polyhedrin promoter. Successful insertion of the polypeptide-encoding sequence will render the polyhedrin gene inactive and produce recombinant virus lacking coat protein. The recombinant viruses may then be used to infect, for example, *S. frugiperda* cells or *Trichoplusia* larvae in which the polypeptide of interest may be expressed (Engelhard, E. K. *et al.* (1994) *Proc. Natl. Acad. Sci.* 91 :3224-3227).

In mammalian host cells, a number of viral-based expression systems are generally available. For example, in cases where an adenovirus is used as an expression vector, sequences encoding a polypeptide of interest may be ligated into an adenovirus transcription/translation complex consisting of the late promoter and tripartite leader sequence. Insertion in a non-essential E1 or E3 region of the viral genome may be used to obtain a viable virus which is capable of expressing the polypeptide in infected host cells (Logan, J. and Shenk, T. (1984) *Proc. Natl. Acad. Sci.* 81:3655-3659). In addition,

transcription enhancers, such as the Rous sarcoma virus (RSV) enhancer, may be used to increase expression in mammalian host cells.

Specific initiation signals may also be used to achieve more efficient translation of sequences encoding a polypeptide of interest. Such signals include the ATG initiation codon and adjacent sequences. In cases where sequences encoding the polypeptide, its initiation codon, and upstream sequences are inserted into the appropriate expression vector, no additional transcriptional or translational control signals may be needed. However, in cases where only coding sequence, or a portion thereof, is inserted, exogenous translational control signals including the ATG initiation codon should be provided. Furthermore, the initiation codon should be in the correct reading frame to ensure translation of the entire insert. Exogenous translational elements and initiation codons may be of various origins, both natural and synthetic. The efficiency of expression may be enhanced by the inclusion of enhancers which are appropriate for the particular cell system which is used, such as those described in the literature (Scharf, D. *et al.* (1994) *Results* *Probl. Cell Differ.* 20:125-162).

In addition, a host cell strain may be chosen for its ability to modulate the expression of the inserted sequences or to process the expressed protein in the desired fashion. Such modifications of the polypeptide include, but are not limited to, acetylation, carboxylation, glycosylation, phosphorylation, lipidation, and acylation. Post-translational processing which cleaves a "prepro" form of the protein may also be used to facilitate correct insertion, folding and/or function. Different host cells such as CHO, HeLa, MDCK, HEK293, and WI38, which have specific cellular machinery and characteristic mechanisms for such post-translational activities, may be chosen to ensure the correct modification and processing of the foreign protein.

For long-term, high-yield production of recombinant proteins, stable expression is generally preferred. For example, cell lines which stably express a polynucleotide of interest may be transformed using expression vectors which may contain viral origins of replication and/or endogenous expression elements and a selectable marker gene on the same or on a separate vector. Following the introduction of the vector, cells

may be allowed to grow for 1-2 days in an enriched media before they are switched to selective media. The purpose of the selectable marker is to confer resistance to selection, and its presence allows growth and recovery of cells which successfully express the introduced sequences. Resistant clones of stably transformed cells may be proliferated
5 using tissue culture techniques appropriate to the cell type.

Any number of selection systems may be used to recover transformed cell lines. These include, but are not limited to, the herpes simplex virus thymidine kinase (Wigler, M. *et al.* (1977) *Cell* 11:223-32) and adenine phosphoribosyltransferase (Lowy, I. *et al.* (1990) *Cell* 22:817-23) genes which can be employed in tk.sup.- or aprt.sup.- cells,
10 respectively. Also, antimetabolite, antibiotic or herbicide resistance can be used as the basis for selection; for example, dhfr which confers resistance to methotrexate (Wigler, M. *et al.* (1980) *Proc. Natl. Acad. Sci.* 77:3567-70); npt, which confers resistance to the aminoglycosides, neomycin and G-418 (Colbere-Garapin, F. *et al.* (1981) *J. Mol. Biol.* 150:1-14); and als or pat, which confer resistance to chlorsulfuron and phosphinotricin
15 acetyltransferase, respectively (Murry, *supra*). Additional selectable genes have been described, for example, trpB, which allows cells to utilize indole in place of tryptophan, or hisD, which allows cells to utilize histinol in place of histidine (Hartman, S. C. and R. C. Mulligan (1988) *Proc. Natl. Acad. Sci.* 85:8047-51). Recently, the use of visible markers has gained popularity with such markers as anthocyanins, beta-glucuronidase and its
20 substrate GUS, and luciferase and its substrate luciferin, being widely used not only to identify transformants, but also to quantify the amount of transient or stable protein expression attributable to a specific vector system (Rhodes, C. A. *et al.* (1995) *Methods Mol. Biol.* 55:121-131).

Although the presence/absence of marker gene expression suggests that the
25 gene of interest is also present, its presence and expression may need to be confirmed. For example, if the sequence encoding a polypeptide is inserted within a marker gene sequence, recombinant cells containing sequences can be identified by the absence of marker gene function. Alternatively, a marker gene can be placed in tandem with a polypeptide-encoding sequence under the control of a single promoter. Expression of the marker gene

in response to induction or selection usually indicates expression of the tandem gene as well.

Alternatively, host cells which contain and express a desired polynucleotide sequence may be identified by a variety of procedures known to those of skill in the art.

- 5 These procedures include, but are not limited to, DNA-DNA or DNA-RNA hybridizations and protein bioassay or immunoassay techniques which include membrane, solution, or chip based technologies for the detection and/or quantification of nucleic acid or protein.

A variety of protocols for detecting and measuring the expression of polynucleotide-encoded products, using either polyclonal or monoclonal antibodies
 10 specific for the product are known in the art. Examples include enzyme-linked immunosorbent assay (ELISA), radioimmunoassay (RIA), and fluorescence activated cell sorting (FACS). A two-site, monoclonal-based immunoassay utilizing monoclonal antibodies reactive to two non-interfering epitopes on a given polypeptide may be preferred for some applications, but a competitive binding assay may also be employed. These and
 15 other assays are described, among other places, in Hampton, R. *et al.* (1990; Serological Methods, a Laboratory Manual, APS Press, St Paul, Minn.) and Maddox, D. E. *et al.* (1983; *J. Exp. Med.* 158:1211-1216).

A wide variety of labels and conjugation techniques are known by those skilled in the art and may be used in various nucleic acid and amino acid assays. Means for
 20 producing labeled hybridization or PCR probes for detecting sequences related to polynucleotides include oligolabeling, nick translation, end-labeling or PCR amplification using a labeled nucleotide. Alternatively, the sequences, or any portions thereof may be cloned into a vector for the production of an mRNA probe. Such vectors are known in the art, are commercially available, and may be used to synthesize RNA probes in vitro by
 25 addition of an appropriate RNA polymerase such as T7, T3, or SP6 and labeled nucleotides. These procedures may be conducted using a variety of commercially available kits. Suitable reporter molecules or labels, which may be used include radionuclides, enzymes, fluorescent, chemiluminescent, or chromogenic agents as well as substrates, cofactors, inhibitors, magnetic particles, and the like.

Host cells transformed with a polynucleotide sequence of interest may be cultured under conditions suitable for the expression and recovery of the protein from cell culture. The protein produced by a recombinant cell may be secreted or contained intracellularly depending on the sequence and/or the vector used. As will be understood by
 5 those of skill in the art, expression vectors containing polynucleotides of the invention may be designed to contain signal sequences which direct secretion of the encoded polypeptide through a prokaryotic or eukaryotic cell membrane. Other recombinant constructions may be used to join sequences encoding a polypeptide of interest to nucleotide sequence encoding a polypeptide domain which will facilitate purification of soluble proteins. Such
 10 purification facilitating domains include, but are not limited to, metal chelating peptides such as histidine-tryptophan modules that allow purification on immobilized metals, protein A domains that allow purification on immobilized immunoglobulin, and the domain utilized in the FLAGS extension/affinity purification system (Immunex Corp., Seattle, Wash.). The inclusion of cleavable linker sequences such as those specific for Factor XA or
 15 enterokinase (Invitrogen, San Diego, Calif.) between the purification domain and the encoded polypeptide may be used to facilitate purification. One such expression vector provides for expression of a fusion protein containing a polypeptide of interest and a nucleic acid encoding 6 histidine residues preceding a thioredoxin or an enterokinase cleavage site. The histidine residues facilitate purification on IMIAC (immobilized metal
 20 ion affinity chromatography) as described in Porath, J. *et al.* (1992, *Prot. Exp. Purif.* 3:263-281) while the enterokinase cleavage site provides a means for purifying the desired polypeptide from the fusion protein. A discussion of vectors which contain fusion proteins is provided in Kroll, D. J. *et al.* (1993; *DNA Cell Biol.* 12:441-453).

In addition to recombinant production methods, polypeptides of the
 25 invention, and fragments thereof, may be produced by direct peptide synthesis using solid-phase techniques (Merrifield J. (1963) *J. Am. Chem. Soc.* 85:2149-2154). Protein synthesis may be performed using manual techniques or by automation. Automated synthesis may be achieved, for example, using Applied Biosystems 431A Peptide Synthesizer (Perkin

Elmer). Alternatively, various fragments may be chemically synthesized separately and combined using chemical methods to produce the full length molecule.

SITE-SPECIFIC MUTAGENESIS

Site-specific mutagenesis is a technique useful in the preparation of individual peptides, or biologically functional equivalent polypeptides, through specific mutagenesis of the underlying polynucleotides that encode them. The technique, well-known to those of skill in the art, further provides a ready ability to prepare and test sequence variants, for example, incorporating one or more of the foregoing considerations, by introducing one or more nucleotide sequence changes into the DNA. Site-specific mutagenesis allows the production of mutants through the use of specific oligonucleotide sequences which encode the DNA sequence of the desired mutation, as well as a sufficient number of adjacent nucleotides, to provide a primer sequence of sufficient size and sequence complexity to form a stable duplex on both sides of the deletion junction being traversed. Mutations may be employed in a selected polynucleotide sequence to improve, alter, decrease, modify, or otherwise change the properties of the polynucleotide itself, and/or alter the properties, activity, composition, stability, or primary sequence of the encoded polypeptide.

In certain embodiments of the present invention, the inventors contemplate the mutagenesis of the disclosed polynucleotide sequences to alter one or more properties of the encoded polypeptide, such as the antigenicity of a polypeptide vaccine. The techniques of site-specific mutagenesis are well-known in the art, and are widely used to create variants of both polypeptides and polynucleotides. For example, site-specific mutagenesis is often used to alter a specific portion of a DNA molecule. In such embodiments, a primer comprising typically about 14 to about 25 nucleotides or so in length is employed, with about 5 to about 10 residues on both sides of the junction of the sequence being altered.

As will be appreciated by those of skill in the art, site-specific mutagenesis techniques have often employed a phage vector that exists in both a single stranded and

double stranded form. Typical vectors useful in site-directed mutagenesis include vectors such as the M13 phage. These phage are readily commercially-available and their use is generally well-known to those skilled in the art. Double-stranded plasmids are also routinely employed in site directed mutagenesis that eliminates the step of transferring the gene of interest from a plasmid to a phage.

In general, site-directed mutagenesis in accordance herewith is performed by first obtaining a single-stranded vector or melting apart of two strands of a double-stranded vector that includes within its sequence a DNA sequence that encodes the desired peptide. An oligonucleotide primer bearing the desired mutated sequence is prepared, generally synthetically. This primer is then annealed with the single-stranded vector, and subjected to DNA polymerizing enzymes such as *E. coli* polymerase I Klenow fragment, in order to complete the synthesis of the mutation-bearing strand. Thus, a heteroduplex is formed wherein one strand encodes the original non-mutated sequence and the second strand bears the desired mutation. This heteroduplex vector is then used to transform appropriate cells, such as *E. coli* cells, and clones are selected which include recombinant vectors bearing the mutated sequence arrangement.

The preparation of sequence variants of the selected peptide-encoding DNA segments using site-directed mutagenesis provides a means of producing potentially useful species and is not meant to be limiting as there are other ways in which sequence variants of peptides and the DNA sequences encoding them may be obtained. For example, recombinant vectors encoding the desired peptide sequence may be treated with mutagenic agents, such as hydroxylamine, to obtain sequence variants. Specific details regarding these methods and protocols are found in the teachings of Maloy *et al.*, 1994; Segal, 1976; Prokop and Bajpai, 1991; Kuby, 1994; and Maniatis *et al.*, 1982, each incorporated herein by reference, for that purpose.

As used herein, the term "oligonucleotide directed mutagenesis procedure" refers to template-dependent processes and vector-mediated propagation which result in an increase in the concentration of a specific nucleic acid molecule relative to its initial concentration, or in an increase in the concentration of a detectable signal, such as

amplification. As used herein, the term "oligonucleotide directed mutagenesis procedure" is intended to refer to a process that involves the template-dependent extension of a primer molecule. The term template dependent process refers to nucleic acid synthesis of an RNA or a DNA molecule wherein the sequence of the newly synthesized strand of nucleic acid is dictated by the well-known rules of complementary base pairing (see, for example, Watson, 1987). Typically, vector mediated methodologies involve the introduction of the nucleic acid fragment into a DNA or RNA vector, the clonal amplification of the vector, and the recovery of the amplified nucleic acid fragment. Examples of such methodologies are provided by U. S. Patent No. 4,237,224, specifically incorporated herein by reference in its entirety.

POLYNUCLEOTIDE AMPLIFICATION TECHNIQUES

A number of template dependent processes are available to amplify the target sequences of interest present in a sample. One of the best known amplification methods is the polymerase chain reaction (PCR™) which is described in detail in U.S. Patent Nos. 4,683,195, 4,683,202 and 4,800,159, each of which is incorporated herein by reference in its entirety. Briefly, in PCR™, two primer sequences are prepared which are complementary to regions on opposite complementary strands of the target sequence. An excess of deoxynucleoside triphosphates is added to a reaction mixture along with a DNA polymerase (e.g., *Taq* polymerase). If the target sequence is present in a sample, the primers will bind to the target and the polymerase will cause the primers to be extended along the target sequence by adding on nucleotides. By raising and lowering the temperature of the reaction mixture, the extended primers will dissociate from the target to form reaction products, excess primers will bind to the target and to the reaction product and the process is repeated. Preferably reverse transcription and PCR™ amplification procedure may be performed in order to quantify the amount of mRNA amplified. Polymerase chain reaction methodologies are well known in the art.

Another method for amplification is the ligase chain reaction (referred to as LCR), disclosed in Eur. Pat. Appl. Publ. No. 320,308 (specifically incorporated herein by

reference in its entirety). In LCR, two complementary probe pairs are prepared, and in the presence of the target sequence, each pair will bind to opposite complementary strands of the target such that they abut. In the presence of a ligase, the two probe pairs will link to form a single unit. By temperature cycling, as in PCR™, bound ligated units dissociate
 5 from the target and then serve as "target sequences" for ligation of excess probe pairs. U.S. Patent No. 4,883,750, incorporated herein by reference in its entirety, describes an alternative method of amplification similar to LCR for binding probe pairs to a target sequence.

Qbeta Replicase, described in PCT Intl. Pat. Appl. Publ. No.
 10 PCT/US87/00880, incorporated herein by reference in its entirety, may also be used as still another amplification method in the present invention. In this method, a replicative sequence of RNA that has a region complementary to that of a target is added to a sample in the presence of an RNA polymerase. The polymerase will copy the replicative sequence that can then be detected.

15 An isothermal amplification method, in which restriction endonucleases and ligases are used to achieve the amplification of target molecules that contain nucleotide 5'-[α-thio]triphosphates in one strand of a restriction site (Walker *et al.*, 1992, incorporated herein by reference in its entirety), may also be useful in the amplification of nucleic acids in the present invention.

20 Strand Displacement Amplification (SDA) is another method of carrying out isothermal amplification of nucleic acids which involves multiple rounds of strand displacement and synthesis, *i.e.* nick translation. A similar method, called Repair Chain Reaction (RCR) is another method of amplification which may be useful in the present invention and is involves annealing several probes throughout a region targeted for
 25 amplification, followed by a repair reaction in which only two of the four bases are present. The other two bases can be added as biotinylated derivatives for easy detection. A similar approach is used in SDA.

Sequences can also be detected using a cyclic probe reaction (CPR). In CPR, a probe having a 3' and 5' sequences of non-target DNA and an internal or "middle"

sequence of the target protein specific RNA is hybridized to DNA which is present in a sample. Upon hybridization, the reaction is treated with RNaseH, and the products of the probe are identified as distinctive products by generating a signal that is released after digestion. The original template is annealed to another cycling probe and the reaction is
 5 repeated. Thus, CPR involves amplifying a signal generated by hybridization of a probe to a target gene specific expressed nucleic acid.

Still other amplification methods described in Great Britain Pat. Appl. No. 2 202 328, and in PCT Intl. Pat. Appl. Publ. No. PCT/US89/01025, each of which is incorporated herein by reference in its entirety, may be used in accordance with the present
 10 invention. In the former application, "modified" primers are used in a PCR-like, template and enzyme dependent synthesis. The primers may be modified by labeling with a capture moiety (*e.g.*, biotin) and/or a detector moiety (*e.g.*, enzyme). In the latter application, an excess of labeled probes is added to a sample. In the presence of the target sequence, the probe binds and is cleaved catalytically. After cleavage, the target sequence is released
 15 intact to be bound by excess probe. Cleavage of the labeled probe signals the presence of the target sequence.

Other nucleic acid amplification procedures include transcription-based amplification systems (TAS) (Kwoh *et al.*, 1989; PCT Intl. Pat. Appl. Publ. No. WO 88/10315, incorporated herein by reference in its entirety), including nucleic acid sequence
 20 based amplification (NASBA) and 3SR. In NASBA, the nucleic acids can be prepared for amplification by standard phenol/chloroform extraction, heat denaturation of a sample, treatment with lysis buffer and minispin columns for isolation of DNA and RNA or guanidinium chloride extraction of RNA. These amplification techniques involve annealing a primer that has sequences specific to the target sequence. Following
 25 polymerization, DNA/RNA hybrids are digested with RNase H while double stranded DNA molecules are heat-denatured again. In either case the single stranded DNA is made fully double stranded by addition of second target-specific primer, followed by polymerization. The double stranded DNA molecules are then multiply transcribed by a polymerase such as T7 or SP6. In an isothermal cyclic reaction, the RNAs are reverse

transcribed into DNA, and transcribed once again with a polymerase such as T7 or SP6. The resulting products, whether truncated or complete, indicate target-specific sequences.

Eur. Pat. Appl. Publ. No. 329,822, incorporated herein by reference in its entirety, disclose a nucleic acid amplification process involving cyclically synthesizing
 5 single-stranded RNA ("ssRNA"), ssDNA, and double-stranded DNA (dsDNA), which may be used in accordance with the present invention. The ssRNA is a first template for a first primer oligonucleotide, which is elongated by reverse transcriptase (RNA-dependent DNA polymerase). The RNA is then removed from resulting DNA:RNA duplex by the action of
 10 ribonuclease H (RNase H, an RNase specific for RNA in a duplex with either DNA or RNA). The resultant ssDNA is a second template for a second primer, which also includes the sequences of an RNA polymerase promoter (exemplified by T7 RNA polymerase) 5' to its homology to its template. This primer is then extended by DNA polymerase (exemplified by the large "Klenow" fragment of *E. coli* DNA polymerase I), resulting as a
 15 double-stranded DNA ("dsDNA") molecule, having a sequence identical to that of the original RNA between the primers and having additionally, at one end, a promoter sequence. This promoter sequence can be used by the appropriate RNA polymerase to make many RNA copies of the DNA. These copies can then re-enter the cycle leading to very swift amplification. With proper choice of enzymes, this amplification can be done
 20 isothermally without addition of enzymes at each cycle. Because of the cyclical nature of this process, the starting sequence can be chosen to be in the form of either DNA or RNA.

PCT Intl. Pat. Appl. Publ. No. WO 89/06700, incorporated herein by reference in its entirety, disclose a nucleic acid sequence amplification scheme based on the hybridization of a promoter/primer sequence to a target single-stranded DNA ("ssDNA") followed by transcription of many RNA copies of the sequence. This scheme is not cyclic;
 25 *i.e.* new templates are not produced from the resultant RNA transcripts. Other amplification methods include "RACE" (Frohman, 1990), and "one-sided PCR" (Ohara, 1989) which are well-known to those of skill in the art.

Methods based on ligation of two (or more) oligonucleotides in the presence of nucleic acid having the sequence of the resulting "di-oligonucleotide", thereby

amplifying the di-oligonucleotide (Wu and Dean, 1996, incorporated herein by reference in its entirety), may also be used in the amplification of DNA sequences of the present invention.

BIOLOGICAL FUNCTIONAL EQUIVALENTS

5 Modification and changes may be made in the structure of the polynucleotides and polypeptides of the present invention and still obtain a functional molecule that encodes a polypeptide with desirable characteristics. As mentioned above, it is often desirable to introduce one or more mutations into a specific polynucleotide sequence. In certain circumstances, the resulting encoded polypeptide sequence is altered
10 by this mutation, or in other cases, the sequence of the polypeptide is unchanged by one or more mutations in the encoding polynucleotide.

When it is desirable to alter the amino acid sequence of a polypeptide to create an equivalent, or even an improved, second-generation molecule, the amino acid changes may be achieved by changing one or more of the codons of the encoding DNA
15 sequence, according to Table 1.

For example, certain amino acids may be substituted for other amino acids in a protein structure without appreciable loss of interactive binding capacity with structures such as, for example, antigen-binding regions of antibodies or binding sites on substrate molecules. Since it is the interactive capacity and nature of a protein that defines
20 that protein's biological functional activity, certain amino acid sequence substitutions can be made in a protein sequence, and, of course, its underlying DNA coding sequence, and nevertheless obtain a protein with like properties. It is thus contemplated by the inventors that various changes may be made in the peptide sequences of the disclosed compositions, or corresponding DNA sequences which encode said peptides without appreciable loss of
25 their biological utility or activity.

TABLE 1

| Amino Acids | | | Codons | | | | | | |
|---------------|-----|---|--------|-----|-----|-----|-----|-----|--|
| Alanine | Ala | A | GCA | GCC | GCG | GCU | | | |
| Cysteine | Cys | C | UGC | UGU | | | | | |
| Aspartic acid | Asp | D | GAC | GAU | | | | | |
| Glutamic acid | Glu | E | GAA | GAG | | | | | |
| Phenylalanine | Phe | F | UUC | UUU | | | | | |
| Glycine | Gly | G | GGA | GGC | GGG | GGU | | | |
| Histidine | His | H | CAC | CAU | | | | | |
| Isoleucine | Ile | I | AUA | AUC | AUU | | | | |
| Lysine | Lys | K | AAA | AAG | | | | | |
| Leucine | Leu | L | UUA | UUG | CUA | CUC | CUG | CUU | |
| Methionine | Met | M | AUG | | | | | | |
| Asparagine | Asn | N | AAC | AAU | | | | | |
| Proline | Pro | P | CCA | CCC | CCG | CCU | | | |
| Glutamine | Gln | Q | CAA | CAG | | | | | |
| Arginine | Arg | R | AGA | AGG | CGA | CGC | CGG | CGU | |
| Serine | Ser | S | AGC | AGU | UCA | UCC | UCG | UCU | |
| Threonine | Thr | T | ACA | ACC | ACG | ACU | | | |
| Valine | Val | V | GUA | GUC | GUG | GUU | | | |
| Tryptophan | Trp | W | UGG | | | | | | |
| Tyrosine | Tyr | Y | UAC | UAU | | | | | |

In making such changes, the hydropathic index of amino acids may be considered. The importance of the hydropathic amino acid index in conferring interactive biologic function on a protein is generally understood in the art (Kyte and Doolittle, 1982, incorporated herein by reference). It is accepted that the relative hydropathic character of the amino acid contributes to the secondary structure of the resultant protein, which in turn defines the interaction of the protein with other molecules, for example, enzymes,

substrates, receptors, DNA, antibodies, antigens, and the like. Each amino acid has been assigned a hydropathic index on the basis of its hydrophobicity and charge characteristics (Kyte and Doolittle, 1982). These values are: isoleucine (+4.5); valine (+4.2); leucine (+3.8); phenylalanine (+2.8); cysteine/cystine (+2.5); methionine (+1.9); alanine (+1.8);
 5 glycine (−0.4); threonine (−0.7); serine (−0.8); tryptophan (−0.9); tyrosine (−1.3); proline (−1.6); histidine (−3.2); glutamate (−3.5); glutamine (−3.5); aspartate (−3.5); asparagine (−3.5); lysine (−3.9); and arginine (−4.5).

It is known in the art that certain amino acids may be substituted by other amino acids having a similar hydropathic index or score and still result in a protein with
 10 similar biological activity, *i.e.* still obtain a biological functionally equivalent protein. In making such changes, the substitution of amino acids whose hydropathic indices are within ± 2 is preferred, those within ± 1 are particularly preferred, and those within ± 0.5 are even more particularly preferred. It is also understood in the art that the substitution of like amino acids can be made effectively on the basis of hydrophilicity. U. S. Patent 4,554,101
 15 (specifically incorporated herein by reference in its entirety), states that the greatest local average hydrophilicity of a protein, as governed by the hydrophilicity of its adjacent amino acids, correlates with a biological property of the protein.

As detailed in U. S. Patent 4,554,101, the following hydrophilicity values have been assigned to amino acid residues: arginine (+3.0); lysine (+3.0); aspartate (+3.0 \pm 1); glutamate (+3.0 \pm 1); serine (+0.3); asparagine (+0.2); glutamine (+0.2); glycine (0);
 20 threonine (−0.4); proline (−0.5 \pm 1); alanine (−0.5); histidine (−0.5); cysteine (−1.0); methionine (−1.3); valine (−1.5); leucine (−1.8); isoleucine (−1.8); tyrosine (−2.3); phenylalanine (−2.5); tryptophan (−3.4). It is understood that an amino acid can be substituted for another having a similar hydrophilicity value and still obtain a biologically
 25 equivalent, and in particular, an immunologically equivalent protein. In such changes, the substitution of amino acids whose hydrophilicity values are within ± 2 is preferred, those within ± 1 are particularly preferred, and those within ± 0.5 are even more particularly preferred.

As outlined above, amino acid substitutions are generally therefore based on the relative similarity of the amino acid side-chain substituents, for example, their hydrophobicity, hydrophilicity, charge, size, and the like. Exemplary substitutions that take various of the foregoing characteristics into consideration are well known to those of skill in the art and include: arginine and lysine; glutamate and aspartate; serine and threonine; glutamine and asparagine; and valine, leucine and isoleucine.

In addition, any polynucleotide may be further modified to increase stability *in vivo*. Possible modifications include, but are not limited to, the addition of flanking sequences at the 5' and/or 3' ends; the use of phosphorothioate or 2' O-methyl rather than phosphodiesterase linkages in the backbone; and/or the inclusion of nontraditional bases such as inosine, queosine and wybutosine, as well as acetyl- methyl-, thio- and other modified forms of adenine, cytidine, guanine, thymine and uridine.

IN VIVO POLYNUCLEOTIDE DELIVERY TECHNIQUES

In additional embodiments, genetic constructs comprising one or more of the polynucleotides of the invention are introduced into cells *in vivo*. This may be achieved using any of a variety of well known approaches, several of which are outlined below for the purpose of illustration.

1. ADENOVIRUS

One of the preferred methods for *in vivo* delivery of one or more nucleic acid sequences involves the use of an adenovirus expression vector. "Adenovirus expression vector" is meant to include those constructs containing adenovirus sequences sufficient to (a) support packaging of the construct and (b) to express a polynucleotide that has been cloned therein in a sense or antisense orientation. Of course, in the context of an antisense construct, expression does not require that the gene product be synthesized.

The expression vector comprises a genetically engineered form of an adenovirus. Knowledge of the genetic organization of adenovirus, a 36 kb, linear, double-stranded DNA virus, allows substitution of large pieces of adenoviral DNA with foreign

Adenovirus is particularly suitable for use as a gene transfer vector because of its mid-sized genome, ease of manipulation, high titer, wide target-cell range and high infectivity. Both ends of the viral genome contain 100-200 base pair inverted repeats (ITRs), which are *cis* elements necessary for viral DNA replication and packaging. The early (E) and late (L) regions of the genome contain different transcription units that are divided by the onset of viral DNA replication. The E1 region (E1A and E1B) encodes proteins responsible for the regulation of transcription of the viral genome and a few cellular genes. The expression of the E2 region (E2A and E2B) results in the synthesis of the proteins for viral DNA replication. These proteins are involved in DNA replication, late gene expression and host cell shut-off (Renan, 1990). The products of the late genes, including the majority of the viral capsid proteins, are expressed only after significant processing of a single primary transcript issued by the major late promoter (MLP). The MLP, (located at 16.8 m.u.) is particularly efficient during the late phase of infection, and all the mRNA's issued from this promoter possess a 5'-tripartite leader (TPL) sequence which makes them preferred mRNA's for translation.

Generation and propagation of the current adenovirus vectors, which are replication deficient, depend on a unique helper cell line, designated 293, which was

transformed from human embryonic kidney cells by Ad5 DNA fragments and constitutively expresses E1 proteins (Graham *et al.*, 1977). Since the E3 region is dispensable from the adenovirus genome (Jones and Shenk, 1978), the current adenovirus vectors, with the help of 293 cells, carry foreign DNA in either the E1, the D3 or both
 5 regions (Graham and Prevec, 1991). In nature, adenovirus can package approximately 105% of the wild-type genome (Ghosh-Choudhury *et al.*, 1987), providing capacity for about 2 extra kB of DNA. Combined with the approximately 5.5 kB of DNA that is replaceable in the E1 and E3 regions, the maximum capacity of the current adenovirus vector is under 7.5 kB, or about 15% of the total length of the vector. More than 80% of
 10 the adenovirus viral genome remains in the vector backbone and is the source of vector-borne cytotoxicity. Also, the replication deficiency of the E1-deleted virus is incomplete. For example, leakage of viral gene expression has been observed with the currently available vectors at high multiplicities of infection (MOI) (Mulligan, 1993).

Helper cell lines may be derived from human cells such as human
 15 embryonic kidney cells, muscle cells, hematopoietic cells or other human embryonic mesenchymal or epithelial cells. Alternatively, the helper cells may be derived from the cells of other mammalian species that are permissive for human adenovirus. Such cells include, *e.g.*, Vero cells or other monkey embryonic mesenchymal or epithelial cells. As stated above, the currently preferred helper cell line is 293.

20 Recently, Racher *et al.* (1995) disclosed improved methods for culturing 293 cells and propagating adenovirus. In one format, natural cell aggregates are grown by inoculating individual cells into 1 liter siliconized spinner flasks (Techne, Cambridge, UK) containing 100-200 ml of medium. Following stirring at 40 rpm, the cell viability is estimated with trypan blue. In another format, Fibra-Cel microcarriers (Bibby Sterlin,
 25 Stone, UK) (5 g/l) is employed as follows. A cell inoculum, resuspended in 5 ml of medium, is added to the carrier (50 ml) in a 250 ml Erlenmeyer flask and left stationary, with occasional agitation, for 1 to 4 h. The medium is then replaced with 50 ml of fresh medium and shaking initiated. For virus production, cells are allowed to grow to about 80% confluence, after which time the medium is replaced (to 25% of the final volume) and

adenovirus added at an MOI of 0.05. Cultures are left stationary overnight, following which the volume is increased to 100% and shaking commenced for another 72 h.

Other than the requirement that the adenovirus vector be replication defective, or at least conditionally defective, the nature of the adenovirus vector is not
 5 believed to be crucial to the successful practice of the invention. The adenovirus may be of any of the 42 different known serotypes or subgroups A-F. Adenovirus type 5 of subgroup C is the preferred starting material in order to obtain a conditional replication-defective adenovirus vector for use in the present invention, since Adenovirus type 5 is a human adenovirus about which a great deal of biochemical and genetic information is known, and
 10 it has historically been used for most constructions employing adenovirus as a vector.

As stated above, the typical vector according to the present invention is replication defective and will not have an adenovirus E1 region. Thus, it will be most convenient to introduce the polynucleotide encoding the gene of interest at the position from which the E1-coding sequences have been removed. However, the position of
 15 insertion of the construct within the adenovirus sequences is not critical to the invention. The polynucleotide encoding the gene of interest may also be inserted in lieu of the deleted E3 region in E3 replacement vectors as described by Karlsson *et al.* (1986) or in the E4 region where a helper cell line or helper virus complements the E4 defect.

Adenovirus is easy to grow and manipulate and exhibits broad host range *in vitro* and *in vivo*. This group of viruses can be obtained in high titers, *e.g.*, 10^9 - 10^{11} plaque-forming units per ml, and they are highly infective. The life cycle of adenovirus does not require integration into the host cell genome. The foreign genes delivered by adenovirus vectors are episomal and, therefore, have low genotoxicity to host cells. No side effects have been reported in studies of vaccination with wild-type adenovirus (Couch *et al.*, 1963;
 25 Top *et al.*, 1971), demonstrating their safety and therapeutic potential as *in vivo* gene transfer vectors.

Adenovirus vectors have been used in eukaryotic gene expression (Levrero *et al.*, 1991; Gomez-Foix *et al.*, 1992) and vaccine development (Grunhaus and Horwitz, 1992; Graham and Prevec, 1992). Recently, animal studies suggested that recombinant

adenovirus could be used for gene therapy (Stratford-Perricaudet and Perricaudet, 1991; Stratford-Perricaudet *et al.*, 1990; Rich *et al.*, 1993). Studies in administering recombinant adenovirus to different tissues include trachea instillation (Rosenfeld *et al.*, 1991; Rosenfeld *et al.*, 1992), muscle injection (Ragot *et al.*, 1993), peripheral intravenous
 5 injections (Herz and Gerard, 1993) and stereotactic inoculation into the brain (Le Gal La Salle *et al.*, 1993).

2. RETROVIRUSES

The retroviruses are a group of single-stranded RNA viruses characterized by an ability to convert their RNA to double-stranded DNA in infected cells by a process of
 10 reverse-transcription (Coffin, 1990). The resulting DNA then stably integrates into cellular chromosomes as a provirus and directs synthesis of viral proteins. The integration results in the retention of the viral gene sequences in the recipient cell and its descendants. The retroviral genome contains three genes, gag, pol, and env that code for capsid proteins, polymerase enzyme, and envelope components, respectively. A sequence found upstream
 15 from the gag gene contains a signal for packaging of the genome into virions. Two long terminal repeat (LTR) sequences are present at the 5' and 3' ends of the viral genome. These contain strong promoter and enhancer sequences and are also required for integration in the host cell genome (Coffin, 1990).

In order to construct a retroviral vector, a nucleic acid encoding one or more
 20 oligonucleotide or polynucleotide sequences of interest is inserted into the viral genome in the place of certain viral sequences to produce a virus that is replication-defective. In order to produce virions, a packaging cell line containing the gag, pol, and env genes but without the LTR and packaging components is constructed (Mann *et al.*, 1983). When a recombinant plasmid containing a cDNA, together with the retroviral LTR and packaging
 25 sequences is introduced into this cell line (by calcium phosphate precipitation for example), the packaging sequence allows the RNA transcript of the recombinant plasmid to be packaged into viral particles, which are then secreted into the culture media (Nicolas and Rubenstein, 1988; Temin, 1986; Mann *et al.*, 1983). The media containing the

recombinant retroviruses is then collected, optionally concentrated, and used for gene transfer. Retroviral vectors are able to infect a broad variety of cell types. However, integration and stable expression require the division of host cells (Paskind *et al.*, 1975).

A novel approach designed to allow specific targeting of retrovirus vectors was recently developed based on the chemical modification of a retrovirus by the chemical addition of lactose residues to the viral envelope. This modification could permit the specific infection of hepatocytes *via* sialoglycoprotein receptors.

A different approach to targeting of recombinant retroviruses was designed in which biotinylated antibodies against a retroviral envelope protein and against a specific cell receptor were used. The antibodies were coupled *via* the biotin components by using streptavidin (Roux *et al.*, 1989). Using antibodies against major histocompatibility complex class I and class II antigens, they demonstrated the infection of a variety of human cells that bore those surface antigens with an ecotropic virus *in vitro* (Roux *et al.*, 1989).

3. ADENO-ASSOCIATED VIRUSES

AAV (Ridgeway, 1988; Hermonat and Muzyczka, 1984) is a parovirus, discovered as a contamination of adenoviral stocks. It is a ubiquitous virus (antibodies are present in 85% of the US human population) that has not been linked to any disease. It is also classified as a dependovirus, because its replications is dependent on the presence of a helper virus, such as adenovirus. Five serotypes have been isolated, of which AAV-2 is the best characterized. AAV has a single-stranded linear DNA that is encapsidated into capsid proteins VP1, VP2 and VP3 to form an icosahedral virion of 20 to 24 nm in diameter (Muzyczka and McLaughlin, 1988).

The AAV DNA is approximately 4.7 kilobases long. It contains two open reading frames and is flanked by two ITRs. There are two major genes in the AAV genome: *rep* and *cap*. The *rep* gene codes for proteins responsible for viral replications, whereas *cap* codes for capsid protein VP1-3. Each ITR forms a T-shaped hairpin structure. These terminal repeats are the only essential *cis* components of the AAV for chromosomal integration. Therefore, the AAV can be used as a vector with all viral coding

sequences removed and replaced by the cassette of genes for delivery. Three viral promoters have been identified and named p5, p19, and p40, according to their map position. Transcription from p5 and p19 results in production of rep proteins, and transcription from p40 produces the capsid proteins (Hermonat and Muzyczka, 1984).

5 There are several factors that prompted researchers to study the possibility of using rAAV as an expression vector. One is that the requirements for delivering a gene to integrate into the host chromosome are surprisingly few. It is necessary to have the 145-bp ITRs, which are only 6% of the AAV genome. This leaves room in the vector to assemble a 4.5-kb DNA insertion. While this carrying capacity may prevent the AAV from
10 delivering large genes, it is amply suited for delivering the antisense constructs of the present invention.

AAV is also a good choice of delivery vehicles due to its safety. There is a relatively complicated rescue mechanism: not only wild type adenovirus but also AAV genes are required to mobilize rAAV. Likewise, AAV is not pathogenic and not associated
15 with any disease. The removal of viral coding sequences minimizes immune reactions to viral gene expression, and therefore, rAAV does not evoke an inflammatory response.

4. OTHER VIRAL VECTORS AS EXPRESSION CONSTRUCTS

Other viral vectors may be employed as expression constructs in the present invention for the delivery of oligonucleotide or polynucleotide sequences to a host cell.
20 Vectors derived from viruses such as vaccinia virus (Ridgeway, 1988; Coupar *et al.*, 1988), lentiviruses, polio viruses and herpes viruses may be employed. They offer several attractive features for various mammalian cells (Friedmann, 1989; Ridgeway, 1988; Coupar *et al.*, 1988; Horwich *et al.*, 1990).

With the recent recognition of defective hepatitis B viruses, new insight was
25 gained into the structure-function relationship of different viral sequences. *In vitro* studies showed that the virus could retain the ability for helper-dependent packaging and reverse transcription despite the deletion of up to 80% of its genome (Horwich *et al.*, 1990). This suggested that large portions of the genome could be replaced with foreign genetic

material. The hepatotropism and persistence (integration) were particularly attractive properties for liver-directed gene transfer. Chang *et al.* (1991) introduced the chloramphenicol acetyltransferase (CAT) gene into duck hepatitis B virus genome in the place of the polymerase, surface, and pre-surface coding sequences. It was cotransfected with wild-type virus into an avian hepatoma cell line. Culture media containing high titers of the recombinant virus were used to infect primary duckling hepatocytes. Stable CAT gene expression was detected for at least 24 days after transfection (Chang *et al.*, 1991).

5. NON-VIRAL VECTORS

In order to effect expression of the oligonucleotide or polynucleotide sequences of the present invention, the expression construct must be delivered into a cell. This delivery may be accomplished *in vitro*, as in laboratory procedures for transforming cells lines, or *in vivo* or *ex vivo*, as in the treatment of certain disease states. As described above, one preferred mechanism for delivery is *via* viral infection where the expression construct is encapsulated in an infectious viral particle.

Once the expression construct has been delivered into the cell the nucleic acid encoding the desired oligonucleotide or polynucleotide sequences may be positioned and expressed at different sites. In certain embodiments, the nucleic acid encoding the construct may be stably integrated into the genome of the cell. This integration may be in the specific location and orientation *via* homologous recombination (gene replacement) or it may be integrated in a random, non-specific location (gene augmentation). In yet further embodiments, the nucleic acid may be stably maintained in the cell as a separate, episomal segment of DNA. Such nucleic acid segments or "episomes" encode sequences sufficient to permit maintenance and replication independent of or in synchronization with the host cell cycle. How the expression construct is delivered to a cell and where in the cell the nucleic acid remains is dependent on the type of expression construct employed.

In certain embodiments of the invention, the expression construct comprising one or more oligonucleotide or polynucleotide sequences may simply consist of naked recombinant DNA or plasmids. Transfer of the construct may be performed by any

of the methods mentioned above which physically or chemically permeabilize the cell membrane. This is particularly applicable for transfer *in vitro* but it may be applied to *in vivo* use as well. Dubensky *et al.* (1984) successfully injected polyomavirus DNA in the form of calcium phosphate precipitates into liver and spleen of adult and newborn mice demonstrating active viral replication and acute infection. Benvenisty and Reshef (1986) also demonstrated that direct intraperitoneal injection of calcium phosphate-precipitated plasmids results in expression of the transfected genes. It is envisioned that DNA encoding a gene of interest may also be transferred in a similar manner *in vivo* and express the gene product.

Another embodiment of the invention for transferring a naked DNA expression construct into cells may involve particle bombardment. This method depends on the ability to accelerate DNA-coated microprojectiles to a high velocity allowing them to pierce cell membranes and enter cells without killing them (Klein *et al.*, 1987). Several devices for accelerating small particles have been developed. One such device relies on a high voltage discharge to generate an electrical current, which in turn provides the motive force (Yang *et al.*, 1990). The microprojectiles used have consisted of biologically inert substances such as tungsten or gold beads.

Selected organs including the liver, skin, and muscle tissue of rats and mice have been bombarded *in vivo* (Yang *et al.*, 1990; Zelenin *et al.*, 1991). This may require surgical exposure of the tissue or cells, to eliminate any intervening tissue between the gun and the target organ, *i.e. ex vivo* treatment. Again, DNA encoding a particular gene may be delivered *via* this method and still be incorporated by the present invention.

ANTISENSE OLIGONUCLEOTIDES

The end result of the flow of genetic information is the synthesis of protein. DNA is transcribed by polymerases into messenger RNA and translated on the ribosome to yield a folded, functional protein. Thus there are several steps along the route where protein synthesis can be inhibited. The native DNA segment coding for a polypeptide described herein, as all such mammalian DNA strands, has two strands: a sense strand and

an antisense strand held together by hydrogen bonding. The messenger RNA coding for polypeptide has the same nucleotide sequence as the sense DNA strand except that the DNA thymidine is replaced by uridine. Thus, synthetic antisense nucleotide sequences will bind to a mRNA and inhibit expression of the protein encoded by that mRNA.

5 The targeting of antisense oligonucleotides to mRNA is thus one mechanism to shut down protein synthesis, and, consequently, represents a powerful and targeted therapeutic approach. For example, the synthesis of polygalacturonase and the muscarine type 2 acetylcholine receptor are inhibited by antisense oligonucleotides directed to their respective mRNA sequences (U. S. Patent 5,739,119 and U. S. Patent 5,759,829, each
10 specifically incorporated herein by reference in its entirety). Further, examples of antisense inhibition have been demonstrated with the nuclear protein cyclin, the multiple drug resistance gene (MDG1), ICAM-1, E-selectin, STK-1, striatal GABA_A receptor and human EGF (Jaskulski *et al.*, 1988; Vasanthakumar and Ahmed, 1989; Peris *et al.*, 1998; U. S. Patent 5,801,154; U. S. Patent 5,789,573; U. S. Patent 5,718,709 and U. S. Patent
15 5,610,288, each specifically incorporated herein by reference in its entirety). Antisense constructs have also been described that inhibit and can be used to treat a variety of abnormal cellular proliferations, *e.g.* cancer (U. S. Patent 5,747,470; U. S. Patent 5,591,317 and U. S. Patent 5,783,683, each specifically incorporated herein by reference in its entirety).

20 Therefore, in exemplary embodiments, the invention provides oligonucleotide sequences that comprise all, or a portion of, any sequence that is capable of specifically binding to polynucleotide sequence described herein, or a complement thereof. In one embodiment, the antisense oligonucleotides comprise DNA or derivatives thereof. In another embodiment, the oligonucleotides comprise RNA or derivatives thereof. In a
25 third embodiment, the oligonucleotides are modified DNAs comprising a phosphorothioated modified backbone. In a fourth embodiment, the oligonucleotide sequences comprise peptide nucleic acids or derivatives thereof. In each case, preferred compositions comprise a sequence region that is complementary, and more preferably

substantially-complementary, and even more preferably, completely complementary to one or more portions of polynucleotides disclosed herein.

Selection of antisense compositions specific for a given gene sequence is based upon analysis of the chosen target sequence (*i.e.* in these illustrative examples the rat and human sequences) and determination of secondary structure, T_m , binding energy, relative stability, and antisense compositions were selected based upon their relative inability to form dimers, hairpins, or other secondary structures that would reduce or prohibit specific binding to the target mRNA in a host cell.

Highly preferred target regions of the mRNA, are those which are at or near the AUG translation initiation codon, and those sequences which were substantially complementary to 5' regions of the mRNA. These secondary structure analyses and target site selection considerations were performed using v.4 of the OLIGO primer analysis software (Rychlik, 1997) and the BLASTN 2.0.5 algorithm software (Altschul *et al.*, 1997).

The use of an antisense delivery method employing a short peptide vector, termed MPG (27 residues), is also contemplated. The MPG peptide contains a hydrophobic domain derived from the fusion sequence of HIV gp41 and a hydrophilic domain from the nuclear localization sequence of SV40 T-antigen (Morris *et al.*, 1997). It has been demonstrated that several molecules of the MPG peptide coat the antisense oligonucleotides and can be delivered into cultured mammalian cells in less than 1 hour with relatively high efficiency (90%). Further, the interaction with MPG strongly increases both the stability of the oligonucleotide to nuclease and the ability to cross the plasma membrane (Morris *et al.*, 1997).

RIBOZYMES

Although proteins traditionally have been used for catalysis of nucleic acids, another class of macromolecules has emerged as useful in this endeavor. Ribozymes are RNA-protein complexes that cleave nucleic acids in a site-specific fashion. Ribozymes have specific catalytic domains that possess endonuclease activity (Kim and Cech, 1987; Gerlach *et al.*, 1987; Forster and Symons, 1987). For example, a large number of

ribozymes accelerate phosphoester transfer reactions with a high degree of specificity, often cleaving only one of several phosphoesters in an oligonucleotide substrate (Cech *et al.*, 1981; Michel and Westhof, 1990; Reinhold-Hurek and Shub, 1992). This specificity has been attributed to the requirement that the substrate bind via specific base-pairing
 5 interactions to the internal guide sequence ("IGS") of the ribozyme prior to chemical reaction.

Ribozyme catalysis has primarily been observed as part of sequence-specific cleavage/ligation reactions involving nucleic acids (Joyce, 1989; Cech *et al.*, 1981). For example, U. S. Patent No. 5,354,855 (specifically incorporated herein by reference) reports
 10 that certain ribozymes can act as endonucleases with a sequence specificity greater than that of known ribonucleases and approaching that of the DNA restriction enzymes. Thus, sequence-specific ribozyme-mediated inhibition of gene expression may be particularly suited to therapeutic applications (Scanlon *et al.*, 1991; Sarver *et al.*, 1990). Recently, it was reported that ribozymes elicited genetic changes in some cells lines to which they were
 15 applied; the altered genes included the oncogenes *H-ras*, *c-fos* and genes of HIV. Most of this work involved the modification of a target mRNA, based on a specific mutant codon that is cleaved by a specific ribozyme.

Six basic varieties of naturally-occurring enzymatic RNAs are known presently. Each can catalyze the hydrolysis of RNA phosphodiester bonds *in trans* (and
 20 thus can cleave other RNA molecules) under physiological conditions. In general, enzymatic nucleic acids act by first binding to a target RNA. Such binding occurs through the target binding portion of a enzymatic nucleic acid which is held in close proximity to an enzymatic portion of the molecule that acts to cleave the target RNA. Thus, the enzymatic nucleic acid first recognizes and then binds a target RNA through complementary base-
 25 pairing, and once bound to the correct site, acts enzymatically to cut the target RNA. Strategic cleavage of such a target RNA will destroy its ability to direct synthesis of an encoded protein. After an enzymatic nucleic acid has bound and cleaved its RNA target, it is released from that RNA to search for another target and can repeatedly bind and cleave new targets.

The enzymatic nature of a ribozyme is advantageous over many technologies, such as antisense technology (where a nucleic acid molecule simply binds to a nucleic acid target to block its translation) since the concentration of ribozyme necessary to affect a therapeutic treatment is lower than that of an antisense oligonucleotide. This advantage reflects the ability of the ribozyme to act enzymatically. Thus, a single ribozyme molecule is able to cleave many molecules of target RNA. In addition, the ribozyme is a highly specific inhibitor, with the specificity of inhibition depending not only on the base pairing mechanism of binding to the target RNA, but also on the mechanism of target RNA cleavage. Single mismatches, or base-substitutions, near the site of cleavage can completely eliminate catalytic activity of a ribozyme. Similar mismatches in antisense molecules do not prevent their action (Woolf *et al.*, 1992). Thus, the specificity of action of a ribozyme is greater than that of an antisense oligonucleotide binding the same RNA site.

The enzymatic nucleic acid molecule may be formed in a hammerhead, hairpin, a hepatitis δ virus, group I intron or RNaseP RNA (in association with an RNA guide sequence) or Neurospora VS RNA motif. Examples of hammerhead motifs are described by Rossi *et al.* (1992). Examples of hairpin motifs are described by Hampel *et al.* (Eur. Pat. Appl. Publ. No. EP 0360257), Hampel and Tritz (1989), Hampel *et al.* (1990) and U. S. Patent 5,631,359 (specifically incorporated herein by reference). An example of the hepatitis δ virus motif is described by Perrotta and Been (1992); an example of the RNaseP motif is described by Guerrier-Takada *et al.* (1983); Neurospora VS RNA ribozyme motif is described by Collins (Saville and Collins, 1990; Saville and Collins, 1991; Collins and Olive, 1993); and an example of the Group I intron is described in (U. S. Patent 4,987,071, specifically incorporated herein by reference). All that is important in an enzymatic nucleic acid molecule of this invention is that it has a specific substrate binding site which is complementary to one or more of the target gene RNA regions, and that it have nucleotide sequences within or surrounding that substrate binding site which impart an RNA cleaving activity to the molecule. Thus the ribozyme constructs need not be limited to specific motifs mentioned herein.

In certain embodiments, it may be important to produce enzymatic cleaving agents which exhibit a high degree of specificity for the RNA of a desired target, such as one of the sequences disclosed herein. The enzymatic nucleic acid molecule is preferably targeted to a highly conserved sequence region of a target mRNA. Such enzymatic nucleic acid molecules can be delivered exogenously to specific cells as required. Alternatively, the ribozymes can be expressed from DNA or RNA vectors that are delivered to specific cells.

Small enzymatic nucleic acid motifs (e.g., of the hammerhead or the hairpin structure) may also be used for exogenous delivery. The simple structure of these molecules increases the ability of the enzymatic nucleic acid to invade targeted regions of the mRNA structure. Alternatively, catalytic RNA molecules can be expressed within cells from eukaryotic promoters (e.g., Scanlon *et al.*, 1991; Kashani-Sabet *et al.*, 1992; Dropulic *et al.*, 1992; Weerasinghe *et al.*, 1991; Ojwang *et al.*, 1992; Chen *et al.*, 1992; Sarver *et al.*, 1990). Those skilled in the art realize that any ribozyme can be expressed in eukaryotic cells from the appropriate DNA vector. The activity of such ribozymes can be augmented by their release from the primary transcript by a second ribozyme (Int. Pat. Appl. Publ. No. WO 93/23569, and Int. Pat. Appl. Publ. No. WO 94/02595, both hereby incorporated by reference; Ohkawa *et al.*, 1992; Taira *et al.*, 1991; and Ventura *et al.*, 1993).

Ribozymes may be added directly, or can be complexed with cationic lipids, lipid complexes, packaged within liposomes, or otherwise delivered to target cells. The RNA or RNA complexes can be locally administered to relevant tissues *ex vivo*, or *in vivo* through injection, aerosol inhalation, infusion pump or stent, with or without their incorporation in biopolymers.

Ribozymes may be designed as described in Int. Pat. Appl. Publ. No. WO 93/23569 and Int. Pat. Appl. Publ. No. WO 94/02595, each specifically incorporated herein by reference) and synthesized to be tested *in vitro* and *in vivo*, as described. Such ribozymes can also be optimized for delivery. While specific examples are provided, those in the art will recognize that equivalent RNA targets in other species can be utilized when necessary.

Hammerhead or hairpin ribozymes may be individually analyzed by computer folding (Jaeger *et al.*, 1989) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core are eliminated from
 5 consideration. Varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 or so bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Ribozymes of the hammerhead or hairpin motif may be designed to anneal to various sites in the mRNA message, and can be chemically synthesized. The method of
 10 synthesis used follows the procedure for normal RNA synthesis as described in Usman *et al.* (1987) and in Scaringe *et al.* (1990) and makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. Average stepwise coupling yields are typically >98%. Hairpin ribozymes may be synthesized in two parts and annealed to reconstruct an active
 15 ribozyme (Chowrira and Burke, 1992). Ribozymes may be modified extensively to enhance stability by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-H (for a review see *e.g.*, Usman and Cedergren, 1992). Ribozymes may be purified by gel electrophoresis using general methods or by high pressure liquid chromatography and resuspended in water.

20 Ribozyme activity can be optimized by altering the length of the ribozyme binding arms, or chemically synthesizing ribozymes with modifications that prevent their degradation by serum ribonucleases (see *e.g.*, Int. Pat. Appl. Publ. No. WO 92/07065; Perrault *et al.*, 1990; Pieken *et al.*, 1991; Usman and Cedergren, 1992; Int. Pat. Appl. Publ. No. WO 93/15187; Int. Pat. Appl. Publ. No. WO 91/03162; Eur. Pat. Appl. Publ. No. 92110298.4; U. S. Patent 5,334,711; and Int. Pat. Appl. Publ. No. WO 94/13688,
 25 which describe various chemical modifications that can be made to the sugar moieties of enzymatic RNA molecules), modifications which enhance their efficacy in cells, and removal of stem II bases to shorten RNA synthesis times and reduce chemical requirements.

Sullivan *et al.* (Int. Pat. Appl. Publ. No. WO 94/02595) describes the general methods for delivery of enzymatic RNA molecules. Ribozymes may be administered to cells by a variety of methods known to those familiar to the art, including, but not restricted to, encapsulation in liposomes, by iontophoresis, or by incorporation into
 5 other vehicles, such as hydrogels, cyclodextrins, biodegradable nanocapsules, and bioadhesive microspheres. For some indications, ribozymes may be directly delivered *ex vivo* to cells or tissues with or without the aforementioned vehicles. Alternatively, the RNA/vehicle combination may be locally delivered by direct inhalation, by direct injection or by use of a catheter, infusion pump or stent. Other routes of delivery include, but are not
 10 limited to, intravascular, intramuscular, subcutaneous or joint injection, aerosol inhalation, oral (tablet or pill form), topical, systemic, ocular, intraperitoneal and/or intrathecal delivery. More detailed descriptions of ribozyme delivery and administration are provided in Int. Pat. Appl. Publ. No. WO 94/02595 and Int. Pat. Appl. Publ. No. WO 93/23569, each specifically incorporated herein by reference.

15 Another means of accumulating high concentrations of a ribozyme(s) within cells is to incorporate the ribozyme-encoding sequences into a DNA expression vector. Transcription of the ribozyme sequences are driven from a promoter for eukaryotic RNA polymerase I (pol I), RNA polymerase II (pol II), or RNA polymerase III (pol III). Transcripts from pol II or pol III promoters will be expressed at high levels in all cells; the
 20 levels of a given pol II promoter in a given cell type will depend on the nature of the gene regulatory sequences (enhancers, silencers, *etc.*) present nearby. Prokaryotic RNA polymerase promoters may also be used, providing that the prokaryotic RNA polymerase enzyme is expressed in the appropriate cells (Elroy-Stein and Moss, 1990; Gao and Huang, 1993; Lieber *et al.*, 1993; Zhou *et al.*, 1990). Ribozymes expressed from such promoters
 25 can function in mammalian cells (*e.g.* Kashani-Saber *et al.*, 1992; Ojwang *et al.*, 1992; Chen *et al.*, 1992; Yu *et al.*, 1993; L'Huillier *et al.*, 1992; Lisiewicz *et al.*, 1993). Such transcription units can be incorporated into a variety of vectors for introduction into mammalian cells, including but not restricted to, plasmid DNA vectors, viral DNA vectors

(such as adenovirus or adeno-associated vectors), or viral RNA vectors (such as retroviral, semliki forest virus, sindbis virus vectors).

Ribozymes may be used as diagnostic tools to examine genetic drift and mutations within diseased cells. They can also be used to assess levels of the target RNA molecule. The close relationship between ribozyme activity and the structure of the target RNA allows the detection of mutations in any region of the molecule which alters the base-pairing and three-dimensional structure of the target RNA. By using multiple ribozymes, one may map nucleotide changes which are important to RNA structure and function *in vitro*, as well as in cells and tissues. Cleavage of target RNAs with ribozymes may be used to inhibit gene expression and define the role (essentially) of specified gene products in the progression of disease. In this manner, other genetic targets may be defined as important mediators of the disease. These studies will lead to better treatment of the disease progression by affording the possibility of combinational therapies (*e.g.*, multiple ribozymes targeted to different genes, ribozymes coupled with known small molecule inhibitors, or intermittent treatment with combinations of ribozymes and/or other chemical or biological molecules). Other *in vitro* uses of ribozymes are well known in the art, and include detection of the presence of mRNA associated with an IL-5 related condition. Such RNA is detected by determining the presence of a cleavage product after treatment with a ribozyme using standard methodology.

20 PEPTIDE NUCLEIC ACIDS

In certain embodiments, the inventors contemplate the use of peptide nucleic acids (PNAs) in the practice of the methods of the invention. PNA is a DNA mimic in which the nucleobases are attached to a pseudopeptide backbone (Good and Nielsen, 1997). PNA is able to be utilized in a number methods that traditionally have used RNA or DNA. Often PNA sequences perform better in techniques than the corresponding RNA or DNA sequences and have utilities that are not inherent to RNA or DNA. A review of PNA including methods of making, characteristics of, and methods of using, is provided by Corey (1997) and is incorporated herein by reference. As such, in certain embodiments,

one may prepare PNA sequences that are complementary to one or more portions of the ACE mRNA sequence, and such PNA compositions may be used to regulate, alter, decrease, or reduce the translation of ACE-specific mRNA, and thereby alter the level of ACE activity in a host cell to which such PNA compositions have been administered.

5 PNA have 2-aminoethyl-glycine linkages replacing the normal phosphodiester backbone of DNA (Nielsen *et al.*, 1991; Hanvey *et al.*, 1992; Hyrup and Nielsen, 1996; Neilsen, 1996). This chemistry has three important consequences: firstly, in contrast to DNA or phosphorothioate oligonucleotides, PNAs are neutral molecules; secondly, PNAs are achiral, which avoids the need to develop a stereoselective synthesis; 10 and thirdly, PNA synthesis uses standard Boc (Dueholm *et al.*, 1994) or Fmoc (Thomson *et al.*, 1995) protocols for solid-phase peptide synthesis, although other methods, including a modified Merrifield method, have been used (Christensen *et al.*, 1995).

PNA monomers or ready-made oligomers are commercially available from PerSeptive Biosystems (Framingham, MA). PNA syntheses by either Boc or Fmoc 15 protocols are straightforward using manual or automated protocols (Norton *et al.*, 1995). The manual protocol lends itself to the production of chemically modified PNAs or the simultaneous synthesis of families of closely related PNAs.

As with peptide synthesis, the success of a particular PNA synthesis will depend on the properties of the chosen sequence. For example, while in theory PNAs can 20 incorporate any combination of nucleotide bases, the presence of adjacent purines can lead to deletions of one or more residues in the product. In expectation of this difficulty, it is suggested that, in producing PNAs with adjacent purines, one should repeat the coupling of residues likely to be added inefficiently. This should be followed by the purification of PNAs by reverse-phase high-pressure liquid chromatography (Norton *et al.*, 1995) 25 providing yields and purity of product similar to those observed during the synthesis of peptides.

Modifications of PNAs for a given application may be accomplished by coupling amino acids during solid-phase synthesis or by attaching compounds that contain a carboxylic acid group to the exposed N-terminal amine. Alternatively, PNAs can be

modified after synthesis by coupling to an introduced lysine or cysteine. The ease with which PNAs can be modified facilitates optimization for better solubility or for specific functional requirements. Once synthesized, the identity of PNAs and their derivatives can be confirmed by mass spectrometry. Several studies have made and utilized modifications of PNAs (Norton *et al.*, 1995; Haaime *et al.*, 1996; Stetsenko *et al.*, 1996; Petersen *et al.*, 1995; Ulmann *et al.*, 1996; Koch *et al.*, 1995; Orum *et al.*, 1995; Footer *et al.*, 1996; Griffith *et al.*, 1995; Kremsky *et al.*, 1996; Pardridge *et al.*, 1995; Boffa *et al.*, 1995; Landsdorp *et al.*, 1996; Gambacorti-Passerini *et al.*, 1996; Armitage *et al.*, 1997; Seeger *et al.*, 1997; Ruskowski *et al.*, 1997). U.S. Patent No. 5,700,922 discusses PNA-DNA-PNA chimeric molecules and their uses in diagnostics, modulating protein in organisms, and treatment of conditions susceptible to therapeutics.

In contrast to DNA and RNA, which contain negatively charged linkages, the PNA backbone is neutral. In spite of this dramatic alteration, PNAs recognize complementary DNA and RNA by Watson-Crick pairing (Egholm *et al.*, 1993), validating the initial modeling by Nielsen *et al.* (1991). PNAs lack 3' to 5' polarity and can bind in either parallel or antiparallel fashion, with the antiparallel mode being preferred (Egholm *et al.*, 1993).

Hybridization of DNA oligonucleotides to DNA and RNA is destabilized by electrostatic repulsion between the negatively charged phosphate backbones of the complementary strands. By contrast, the absence of charge repulsion in PNA-DNA or PNA-RNA duplexes increases the melting temperature (T_m) and reduces the dependence of T_m on the concentration of mono- or divalent cations (Nielsen *et al.*, 1991). The enhanced rate and affinity of hybridization are significant because they are responsible for the surprising ability of PNAs to perform strand invasion of complementary sequences within relaxed double-stranded DNA. In addition, the efficient hybridization at inverted repeats suggests that PNAs can recognize secondary structure effectively within double-stranded DNA. Enhanced recognition also occurs with PNAs immobilized on surfaces, and Wang *et al.* have shown that support-bound PNAs can be used to detect hybridization events (Wang *et al.*, 1996).

One might expect that tight binding of PNAs to complementary sequences would also increase binding to similar (but not identical) sequences, reducing the sequence specificity of PNA recognition. As with DNA hybridization, however, selective recognition can be achieved by balancing oligomer length and incubation temperature.

- 5 Moreover, selective hybridization of PNAs is encouraged by PNA-DNA hybridization being less tolerant of base mismatches than DNA-DNA hybridization. For example, a single mismatch within a 16 bp PNA-DNA duplex can reduce the T_m by up to 15°C (Egholm *et al.*, 1993). This high level of discrimination has allowed the development of several PNA-based strategies for the analysis of point mutations (Wang *et al.*, 1996; Carlsson *et al.*, 1996; Thiede *et al.*, 1996; Webb and Hurskainen, 1996; Perry-O'Keefe *et al.*, 1996).

- High-affinity binding provides clear advantages for molecular recognition and the development of new applications for PNAs. For example, 11-13 nucleotide PNAs inhibit the activity of telomerase, a ribonucleo-protein that extends telomere ends using an essential RNA template, while the analogous DNA oligomers do not (Norton *et al.*, 1996).

- Neutral PNAs are more hydrophobic than analogous DNA oligomers, and this can lead to difficulty solubilizing them at neutral pH, especially if the PNAs have a high purine content or if they have the potential to form secondary structures. Their solubility can be enhanced by attaching one or more positive charges to the PNA termini (Nielsen *et al.*, 1991).

- Findings by Allfrey and colleagues suggest that strand invasion will occur spontaneously at sequences within chromosomal DNA (Boffa *et al.*, 1995; Boffa *et al.*, 1996). These studies targeted PNAs to triplet repeats of the nucleotides CAG and used this recognition to purify transcriptionally active DNA (Boffa *et al.*, 1995) and to inhibit transcription (Boffa *et al.*, 1996). This result suggests that if PNAs can be delivered within cells then they will have the potential to be general sequence-specific regulators of gene expression. Studies and reviews concerning the use of PNAs as antisense and anti-gene agents include Nielsen *et al.* (1993b), Hanvey *et al.* (1992), and Good and Nielsen (1997).

Koppelhus *et al.* (1997) have used PNAs to inhibit HIV-1 inverse transcription, showing that PNAs may be used for antiviral therapies.

Methods of characterizing the antisense binding properties of PNAs are discussed in Rose (1993) and Jensen *et al.* (1997). Rose uses capillary gel electrophoresis to determine binding of PNAs to their complementary oligonucleotide, measuring the relative binding kinetics and stoichiometry. Similar types of measurements were made by Jensen *et al.* using BIAcore™ technology.

Other applications of PNAs include use in DNA strand invasion (Nielsen *et al.*, 1991), antisense inhibition (Hanvey *et al.*, 1992), mutational analysis (Orum *et al.*, 1993), enhancers of transcription (Mollegaard *et al.*, 1994), nucleic acid purification (Orum *et al.*, 1995), isolation of transcriptionally active genes (Boffa *et al.*, 1995), blocking of transcription factor binding (Vickers *et al.*, 1995), genome cleavage (Veselkov *et al.*, 1996), biosensors (Wang *et al.*, 1996), *in situ* hybridization (Thisted *et al.*, 1996), and in an alternative to Southern blotting (Perry-O'Keefe, 1996).

15 POLYPEPTIDE COMPOSITIONS

The present invention, in other aspects, provides polypeptide compositions. Generally, a polypeptide of the invention will be an isolated polypeptide (or an epitope, variant, or active fragment thereof) derived from a mammalian species. Preferably, the polypeptide is encoded by a polynucleotide sequence disclosed herein or a sequence which hybridizes under moderately stringent conditions to a polynucleotide sequence disclosed herein. Alternatively, the polypeptide may be defined as a polypeptide which comprises a contiguous amino acid sequence from an amino acid sequence disclosed herein, or which polypeptide comprises an entire amino acid sequence disclosed herein.

In the present invention, a polypeptide composition is also understood to comprise one or more polypeptides that are immunologically reactive with antibodies generated against a polypeptide of the invention, particularly a polypeptide having the amino acid sequence disclosed in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809,

1670-1675, or to active fragments, or to variants or biological functional equivalents thereof.

Likewise, a polypeptide composition of the present invention is understood to comprise one or more polypeptides that are capable of eliciting antibodies that are immunologically reactive with one or more polypeptides encoded by one or more contiguous nucleic acid sequences contained in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or to active fragments, or to variants thereof, or to one or more nucleic acid sequences which hybridize to one or more of these sequences under conditions of moderate to high stringency. Particularly illustrative polypeptides include the amino acid sequences disclosed in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809, 827 and 1670-1675..

As used herein, an active fragment of a polypeptide includes a whole or a portion of a polypeptide which is modified by conventional techniques, *e.g.*, mutagenesis, or by addition, deletion, or substitution, but which active fragment exhibits substantially the same structure function, antigenicity, etc., as a polypeptide as described herein.

In certain illustrative embodiments, the polypeptides of the invention will comprise at least an immunogenic portion of a lung tumor protein or a variant thereof, as described herein. As noted above, a "lung tumor protein" is a protein that is expressed by lung tumor cells. Proteins that are lung tumor proteins also react detectably within an immunoassay (such as an ELISA) with antisera from a patient with lung cancer. Polypeptides as described herein may be of any length. Additional sequences derived from the native protein and/or heterologous sequences may be present, and such sequences may (but need not) possess further immunogenic or antigenic properties.

An "immunogenic portion," as used herein is a portion of a protein that is recognized (*i.e.*, specifically bound) by a B-cell and/or T-cell surface antigen receptor. Such immunogenic portions generally comprise at least 5 amino acid residues, more preferably at least 10, and still more preferably at least 20 amino acid residues of a lung tumor protein or a variant thereof. Certain preferred immunogenic portions include peptides in which an N-terminal leader sequence and/or transmembrane domain have been deleted. Other preferred immunogenic portions may contain a small N- and/or C-terminal deletion (*e.g.*, 1-30 amino acids, preferably 5-15 amino acids), relative to the mature protein.

Immunogenic portions may generally be identified using well known techniques, such as those summarized in Paul, *Fundamental Immunology*, 3rd ed., 243-247 (Raven Press, 1993) and references cited therein. Such techniques include screening polypeptides for the ability to react with antigen-specific antibodies, antisera and/or T-cell lines or clones. As used herein, antisera and antibodies are "antigen-specific" if they specifically bind to an antigen (*i.e.*, they react with the protein in an ELISA or other immunoassay, and do not react detectably with unrelated proteins). Such antisera and antibodies may be prepared as described herein, and using well known techniques. An immunogenic portion of a native lung tumor protein is a portion that reacts with such antisera and/or T-cells at a level that is not substantially less than the reactivity of the full length polypeptide (*e.g.*, in an ELISA and/or T-cell reactivity assay). Such immunogenic portions may react within such assays at a level that is similar to or greater than the reactivity of the full length polypeptide. Such screens may generally be performed using methods well known to those of ordinary skill in the art, such as those described in Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. For example, a polypeptide may be immobilized on a solid support and contacted with patient sera to allow binding of antibodies within the sera to the immobilized polypeptide. Unbound sera may then be removed and bound antibodies detected using, for example, ¹²⁵I-labeled Protein A.

As noted above, a composition may comprise a variant of a native lung tumor protein. A polypeptide "variant," as used herein, is a polypeptide that differs from a native lung tumor protein in one or more substitutions, deletions, additions and/or insertions, such that the immunogenicity of the polypeptide is not substantially diminished.

5 In other words, the ability of a variant to react with antigen-specific antisera may be enhanced or unchanged, relative to the native protein, or may be diminished by less than 50%, and preferably less than 20%, relative to the native protein. Such variants may generally be identified by modifying one of the above polypeptide sequences and evaluating the reactivity of the modified polypeptide with antigen-specific antibodies or
10 antisera as described herein. Preferred variants include those in which one or more portions, such as an N-terminal leader sequence or transmembrane domain, have been removed. Other preferred variants include variants in which a small portion (*e.g.*, 1-30 amino acids, preferably 5-15 amino acids) has been removed from the N- and/or C-terminal of the mature protein.

15 Polypeptide variants encompassed by the present invention include those exhibiting at least about 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% or more identity (determined as described above) to the polypeptides disclosed herein.

Preferably, a variant contains conservative substitutions. A "conservative
20 substitution" is one in which an amino acid is substituted for another amino acid that has similar properties, such that one skilled in the art of peptide chemistry would expect the secondary structure and hydrophobic nature of the polypeptide to be substantially unchanged. Amino acid substitutions may generally be made on the basis of similarity in polarity, charge, solubility, hydrophobicity, hydrophilicity and/or the amphipathic nature of
25 the residues. For example, negatively charged amino acids include aspartic acid and glutamic acid; positively charged amino acids include lysine and arginine; and amino acids with uncharged polar head groups having similar hydrophilicity values include leucine, isoleucine and valine; glycine and alanine; asparagine and glutamine; and serine, threonine, phenylalanine and tyrosine. Other groups of amino acids that may represent conservative

changes include: (1) ala, pro, gly, glu, asp, gln, asn, ser, thr; (2) cys, ser, tyr, thr; (3) val, ile, leu, met, ala, phe; (4) lys, arg, his; and (5) phe, tyr, trp, his. A variant may also, or alternatively, contain nonconservative changes. In a preferred embodiment, variant polypeptides differ from a native sequence by substitution, deletion or addition of five
 5 amino acids or fewer. Variants may also (or alternatively) be modified by, for example, the deletion or addition of amino acids that have minimal influence on the immunogenicity, secondary structure and hydropathic nature of the polypeptide.

As noted above, polypeptides may comprise a signal (or leader) sequence at the N-terminal end of the protein, which co-translationally or post-translationally directs
 10 transfer of the protein. The polypeptide may also be conjugated to a linker or other sequence for ease of synthesis, purification or identification of the polypeptide (*e.g.*, poly-His), or to enhance binding of the polypeptide to a solid support. For example, a polypeptide may be conjugated to an immunoglobulin Fc region.

Polypeptides may be prepared using any of a variety of well known
 15 techniques. Recombinant polypeptides encoded by DNA sequences as described above may be readily prepared from the DNA sequences using any of a variety of expression vectors known to those of ordinary skill in the art. Expression may be achieved in any appropriate host cell that has been transformed or transfected with an expression vector containing a DNA molecule that encodes a recombinant polypeptide. Suitable host cells
 20 include prokaryotes, yeast, and higher eukaryotic cells, such as mammalian cells and plant cells. Preferably, the host cells employed are *E. coli*, yeast or a mammalian cell line such as COS or CHO. Supernatants from suitable host/vector systems which secrete recombinant protein or polypeptide into culture media may be first concentrated using a commercially available filter. Following concentration, the concentrate may be applied to
 25 a suitable purification matrix such as an affinity matrix or an ion exchange resin. Finally, one or more reverse phase HPLC steps can be employed to further purify a recombinant polypeptide.

Portions and other variants having less than about 100 amino acids, and generally less than about 50 amino acids, may also be generated by synthetic means, using

techniques well known to those of ordinary skill in the art. For example, such polypeptides may be synthesized using any of the commercially available solid-phase techniques, such as the Merrifield solid-phase synthesis method, where amino acids are sequentially added to a growing amino acid chain. *See* Merrifield, *J. Am. Chem. Soc.* 85:2149-2146, 1963.

- 5 Equipment for automated synthesis of polypeptides is commercially available from suppliers such as Perkin Elmer/Applied BioSystems Division (Foster City, CA), and may be operated according to the manufacturer's instructions.

Within certain specific embodiments, a polypeptide may be a fusion protein that comprises multiple polypeptides as described herein, or that comprises at least one
 10 polypeptide as described herein and an unrelated sequence, such as a known tumor protein. A fusion partner may, for example, assist in providing T helper epitopes (an immunological fusion partner), preferably T helper epitopes recognized by humans, or may assist in expressing the protein (an expression enhancer) at higher yields than the native recombinant protein. Certain preferred fusion partners are both immunological and
 15 expression enhancing fusion partners. Other fusion partners may be selected so as to increase the solubility of the protein or to enable the protein to be targeted to desired intracellular compartments. Still further fusion partners include affinity tags, which facilitate purification of the protein.

Fusion proteins may generally be prepared using standard techniques,
 20 including chemical conjugation. Preferably, a fusion protein is expressed as a recombinant protein, allowing the production of increased levels, relative to a non-fused protein, in an expression system. Briefly, DNA sequences encoding the polypeptide components may be assembled separately, and ligated into an appropriate expression vector. The 3' end of the DNA sequence encoding one polypeptide component is ligated, with or without a peptide
 25 linker, to the 5' end of a DNA sequence encoding the second polypeptide component so that the reading frames of the sequences are in phase. This permits translation into a single fusion protein that retains the biological activity of both component polypeptides.

A peptide linker sequence may be employed to separate the first and second polypeptide components by a distance sufficient to ensure that each polypeptide folds into

its secondary and tertiary structures. Such a peptide linker sequence is incorporated into the fusion protein using standard techniques well known in the art. Suitable peptide linker sequences may be chosen based on the following factors: (1) their ability to adopt a flexible extended conformation; (2) their inability to adopt a secondary structure that could interact with functional epitopes on the first and second polypeptides; and (3) the lack of hydrophobic or charged residues that might react with the polypeptide functional epitopes. Preferred peptide linker sequences contain Gly, Asn and Ser residues. Other near neutral amino acids, such as Thr and Ala may also be used in the linker sequence. Amino acid sequences which may be usefully employed as linkers include those disclosed in Maratea *et al.*, *Gene* 40:39-46, 1985; Murphy *et al.*, *Proc. Natl. Acad. Sci. USA* 83:8258-8262, 1986; U.S. Patent No. 4,935,233 and U.S. Patent No. 4,751,180. The linker sequence may generally be from 1 to about 50 amino acids in length. Linker sequences are not required when the first and second polypeptides have non-essential N-terminal amino acid regions that can be used to separate the functional domains and prevent steric interference.

The ligated DNA sequences are operably linked to suitable transcriptional or translational regulatory elements. The regulatory elements responsible for expression of DNA are located only 5' to the DNA sequence encoding the first polypeptides. Similarly, stop codons required to end translation and transcription termination signals are only present 3' to the DNA sequence encoding the second polypeptide.

Fusion proteins are also provided. Such proteins comprise a polypeptide as described herein together with an unrelated immunogenic protein. Preferably the immunogenic protein is capable of eliciting a recall response. Examples of such proteins include tetanus, tuberculosis and hepatitis proteins (*see*, for example, Stoute *et al.* *New Engl. J. Med.*, 336:86-91, 1997).

Within preferred embodiments, an immunological fusion partner is derived from protein D, a surface protein of the gram-negative bacterium *Haemophilus influenza* B (WO 91/18926). Preferably, a protein D derivative comprises approximately the first third of the protein (*e.g.*, the first N-terminal 100-110 amino acids), and a protein D derivative may be lipidated. Within certain preferred embodiments, the first 109 residues of a

Lipoprotein D fusion partner is included on the N-terminus to provide the polypeptide with additional exogenous T-cell epitopes and to increase the expression level in *E. coli* (thus functioning as an expression enhancer). The lipid tail ensures optimal presentation of the antigen to antigen presenting cells. Other fusion partners include the non-structural protein from influenzae virus, NS1 (hemagglutinin). Typically, the N-terminal 81 amino acids are used, although different fragments that include T-helper epitopes may be used.

In another embodiment, the immunological fusion partner is the protein known as LYTA, or a portion thereof (preferably a C-terminal portion). LYTA is derived from *Streptococcus pneumoniae*, which synthesizes an N-acetyl-L-alanine amidase known as amidase LYTA (encoded by the *LytA* gene; *Gene* 43:265-292, 1986). LYTA is an autolysin that specifically degrades certain bonds in the peptidoglycan backbone. The C-terminal domain of the LYTA protein is responsible for the affinity to the choline or to some choline analogues such as DEAE. This property has been exploited for the development of *E. coli* C-LYTA expressing plasmids useful for expression of fusion proteins. Purification of hybrid proteins containing the C-LYTA fragment at the amino terminus has been described (*see Biotechnology* 10:795-798, 1992). Within a preferred embodiment, a repeat portion of LYTA may be incorporated into a fusion protein. A repeat portion is found in the C-terminal region starting at residue 178. A particularly preferred repeat portion incorporates residues 188-305.

In general, polypeptides (including fusion proteins) and polynucleotides as described herein are isolated. An "isolated" polypeptide or polynucleotide is one that is removed from its original environment. For example, a naturally-occurring protein is isolated if it is separated from some or all of the coexisting materials in the natural system. Preferably, such polypeptides are at least about 90% pure, more preferably at least about 95% pure and most preferably at least about 99% pure. A polynucleotide is considered to be isolated if, for example, it is cloned into a vector that is not a part of the natural environment.

BINDING AGENTS

The present invention further provides agents, such as antibodies and antigen-binding fragments thereof, that specifically bind to a lung tumor protein. As used herein, an antibody, or antigen-binding fragment thereof, is said to "specifically bind" to a lung tumor protein if it reacts at a detectable level (within, for example, an ELISA) with a lung tumor protein, and does not react detectably with unrelated proteins under similar conditions. As used herein, "binding" refers to a noncovalent association between two separate molecules such that a complex is formed. The ability to bind may be evaluated by, for example, determining a binding constant for the formation of the complex. The binding constant is the value obtained when the concentration of the complex is divided by the product of the component concentrations. In general, two compounds are said to "bind," in the context of the present invention, when the binding constant for complex formation exceeds about 10^3 L/mol. The binding constant may be determined using methods well known in the art.

Binding agents may be further capable of differentiating between patients with and without a cancer, such as lung cancer, using the representative assays provided herein. In other words, antibodies or other binding agents that bind to a lung tumor protein will generate a signal indicating the presence of a cancer in at least about 20% of patients with the disease, and will generate a negative signal indicating the absence of the disease in at least about 90% of individuals without the cancer. To determine whether a binding agent satisfies this requirement, biological samples (*e.g.*, blood, sera, sputum, urine and/or tumor biopsies) from patients with and without a cancer (as determined using standard clinical tests) may be assayed as described herein for the presence of polypeptides that bind to the binding agent. It will be apparent that a statistically significant number of samples with and without the disease should be assayed. Each binding agent should satisfy the above criteria; however, those of ordinary skill in the art will recognize that binding agents may be used in combination to improve sensitivity.

Any agent that satisfies the above requirements may be a binding agent. For example, a binding agent may be a ribosome, with or without a peptide component, an

RNA molecule or a polypeptide. In a preferred embodiment, a binding agent is an antibody or an antigen-binding fragment thereof. Antibodies may be prepared by any of a variety of techniques known to those of ordinary skill in the art. See, e.g., Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. In
5 general, antibodies can be produced by cell culture techniques, including the generation of monoclonal antibodies as described herein, or via transfection of antibody genes into suitable bacterial or mammalian cell hosts, in order to allow for the production of recombinant antibodies. In one technique, an immunogen comprising the polypeptide is initially injected into any of a wide variety of mammals (e.g., mice, rats, rabbits, sheep or
10 goats). In this step, the polypeptides of this invention may serve as the immunogen without modification. Alternatively, particularly for relatively short polypeptides, a superior immune response may be elicited if the polypeptide is joined to a carrier protein, such as bovine serum albumin or keyhole limpet hemocyanin. The immunogen is injected into the animal host, preferably according to a predetermined schedule incorporating one or more
15 booster immunizations, and the animals are bled periodically. Polyclonal antibodies specific for the polypeptide may then be purified from such antisera by, for example, affinity chromatography using the polypeptide coupled to a suitable solid support.

Monoclonal antibodies specific for an antigenic polypeptide of interest may be prepared, for example, using the technique of Kohler and Milstein, *Eur. J. Immunol.*
20 6:511-519, 1976, and improvements thereto. Briefly, these methods involve the preparation of immortal cell lines capable of producing antibodies having the desired specificity (i.e., reactivity with the polypeptide of interest). Such cell lines may be produced, for example, from spleen cells obtained from an animal immunized as described above. The spleen cells are then immortalized by, for example, fusion with a myeloma cell
25 fusion partner, preferably one that is syngeneic with the immunized animal. A variety of fusion techniques may be employed. For example, the spleen cells and myeloma cells may be combined with a nonionic detergent for a few minutes and then plated at low density on a selective medium that supports the growth of hybrid cells, but not myeloma cells. A preferred selection technique uses HAT (hypoxanthine, aminopterin, thymidine) selection.

After a sufficient time, usually about 1 to 2 weeks, colonies of hybrids are observed. Single colonies are selected and their culture supernatants tested for binding activity against the polypeptide. Hybridomas having high reactivity and specificity are preferred.

Monoclonal antibodies may be isolated from the supernatants of growing hybridoma colonies. In addition, various techniques may be employed to enhance the yield, such as injection of the hybridoma cell line into the peritoneal cavity of a suitable vertebrate host, such as a mouse. Monoclonal antibodies may then be harvested from the ascites fluid or the blood. Contaminants may be removed from the antibodies by conventional techniques, such as chromatography, gel filtration, precipitation, and extraction. The polypeptides of this invention may be used in the purification process in, for example, an affinity chromatography step.

Within certain embodiments, the use of antigen-binding fragments of antibodies may be preferred. Such fragments include Fab fragments, which may be prepared using standard techniques. Briefly, immunoglobulins may be purified from rabbit serum by affinity chromatography on Protein A bead columns (Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988) and digested by papain to yield Fab and Fc fragments. The Fab and Fc fragments may be separated by affinity chromatography on protein A bead columns.

Monoclonal antibodies of the present invention may be coupled to one or more therapeutic agents. Suitable agents in this regard include radionuclides, differentiation inducers, drugs, toxins, and derivatives thereof. Preferred radionuclides include ^{90}Y , ^{123}I , ^{125}I , ^{131}I , ^{186}Re , ^{188}Re , ^{211}At , and ^{212}Bi . Preferred drugs include methotrexate, and pyrimidine and purine analogs. Preferred differentiation inducers include phorbol esters and butyric acid. Preferred toxins include ricin, abrin, diphtheria toxin, cholera toxin, gelonin, *Pseudomonas* exotoxin, *Shigella* toxin, and pokeweed antiviral protein.

A therapeutic agent may be coupled (*e.g.*, covalently bonded) to a suitable monoclonal antibody either directly or indirectly (*e.g.*, via a linker group). A direct reaction between an agent and an antibody is possible when each possesses a substituent

capable of reacting with the other. For example, a nucleophilic group, such as an amino or sulfhydryl group, on one may be capable of reacting with a carbonyl-containing group, such as an anhydride or an acid halide, or with an alkyl group containing a good leaving group (*e.g.*, a halide) on the other.

5 Alternatively, it may be desirable to couple a therapeutic agent and an antibody via a linker group. A linker group can function as a spacer to distance an antibody from an agent in order to avoid interference with binding capabilities. A linker group can also serve to increase the chemical reactivity of a substituent on an agent or an antibody, and thus increase the coupling efficiency. An increase in chemical reactivity
10 may also facilitate the use of agents, or functional groups on agents, which otherwise would not be possible.

 It will be evident to those skilled in the art that a variety of bifunctional or polyfunctional reagents, both homo- and hetero-functional (such as those described in the catalog of the Pierce Chemical Co., Rockford, IL), may be employed as the linker group.
15 Coupling may be effected, for example, through amino groups, carboxyl groups, sulfhydryl groups or oxidized carbohydrate residues. There are numerous references describing such methodology, *e.g.*, U.S. Patent No. 4,671,958, to Rodwell *et al.*

 Where a therapeutic agent is more potent when free from the antibody portion of the immunoconjugates of the present invention, it may be desirable to use a
20 linker group which is cleavable during or upon internalization into a cell. A number of different cleavable linker groups have been described. The mechanisms for the intracellular release of an agent from these linker groups include cleavage by reduction of a disulfide bond (*e.g.*, U.S. Patent No. 4,489,710, to Spitler), by irradiation of a photolabile bond (*e.g.*, U.S. Patent No. 4,625,014, to Senter *et al.*), by hydrolysis of derivatized amino
25 acid side chains (*e.g.*, U.S. Patent No. 4,638,045, to Kohn *et al.*), by serum complement-mediated hydrolysis (*e.g.*, U.S. Patent No. 4,671,958, to Rodwell *et al.*), and acid-catalyzed hydrolysis (*e.g.*, U.S. Patent No. 4,569,789, to Blattler *et al.*).

 It may be desirable to couple more than one agent to an antibody. In one embodiment, multiple molecules of an agent are coupled to one antibody molecule. In

another embodiment, more than one type of agent may be coupled to one antibody. Regardless of the particular embodiment, immunoconjugates with more than one agent may be prepared in a variety of ways. For example, more than one agent may be coupled directly to an antibody molecule, or linkers that provide multiple sites for attachment can
 5 be used. Alternatively, a carrier can be used.

A carrier may bear the agents in a variety of ways, including covalent bonding either directly or via a linker group. Suitable carriers include proteins such as albumins (*e.g.*, U.S. Patent No. 4,507,234, to Kato *et al.*), peptides and polysaccharides such as aminodextran (*e.g.*, U.S. Patent No. 4,699,784, to Shih *et al.*). A carrier may also
 10 bear an agent by noncovalent bonding or by encapsulation, such as within a liposome vesicle (*e.g.*, U.S. Patent Nos. 4,429,008 and 4,873,088). Carriers specific for radionuclide agents include radiohalogenated small molecules and chelating compounds. For example, U.S. Patent No. 4,735,792 discloses representative radiohalogenated small molecules and their synthesis. A radionuclide chelate may be formed from chelating compounds that
 15 include those containing nitrogen and sulfur atoms as the donor atoms for binding the metal, or metal oxide, radionuclide. For example, U.S. Patent No. 4,673,562, to Davison *et al.* discloses representative chelating compounds and their synthesis.

A variety of routes of administration for the antibodies and immunoconjugates may be used. Typically, administration will be intravenous,
 20 intramuscular, subcutaneous or in the bed of a resected tumor. It will be evident that the precise dose of the antibody/immunoconjugate will vary depending upon the antibody used, the antigen density on the tumor, and the rate of clearance of the antibody.

T CELLS

Immunotherapeutic compositions may also, or alternatively, comprise T
 25 cells specific for a lung tumor protein. Such cells may generally be prepared *in vitro* or *ex vivo*, using standard procedures. For example, T cells may be isolated from bone marrow, peripheral blood, or a fraction of bone marrow or peripheral blood of a patient, using a commercially available cell separation system, such as the Isolex™ System, available from

Nexell Therapeutics, Inc. (Irvine, CA; see also U.S. Patent No. 5,240,856; U.S. Patent No. 5,215,926; WO 89/06280; WO 91/16116 and WO 92/07243). Alternatively, T cells may be derived from related or unrelated humans, non-human mammals, cell lines or cultures.

5 T cells may be stimulated with a lung tumor polypeptide, polynucleotide encoding a lung tumor polypeptide and/or an antigen presenting cell (APC) that expresses such a polypeptide. Such stimulation is performed under conditions and for a time sufficient to permit the generation of T cells that are specific for the polypeptide. Preferably, a lung tumor polypeptide or polynucleotide is present within a delivery vehicle, such as a microsphere, to facilitate the generation of specific T cells.

10 T cells are considered to be specific for a lung tumor polypeptide if the T cells specifically proliferate, secrete cytokines or kill target cells coated with the polypeptide or expressing a gene encoding the polypeptide. T cell specificity may be evaluated using any of a variety of standard techniques. For example, within a chromium release assay or proliferation assay, a stimulation index of more than two fold increase in
15 lysis and/or proliferation, compared to negative controls, indicates T cell specificity. Such assays may be performed, for example, as described in Chen *et al.*, *Cancer Res.* 54:1065-1070, 1994. Alternatively, detection of the proliferation of T cells may be accomplished by a variety of known techniques. For example, T cell proliferation can be detected by measuring an increased rate of DNA synthesis (*e.g.*, by pulse-labeling cultures of T cells
20 with tritiated thymidine and measuring the amount of tritiated thymidine incorporated into DNA). Contact with a lung tumor polypeptide (100 ng/ml - 100 µg/ml, preferably 200 ng/ml - 25 µg/ml) for 3 - 7 days should result in at least a two fold increase in proliferation of the T cells. Contact as described above for 2-3 hours should result in activation of the T cells, as measured using standard cytokine assays in which a two fold increase in the level
25 of cytokine release (*e.g.*, TNF or IFN-γ) is indicative of T cell activation (*see* Coligan *et al.*, *Current Protocols in Immunology*, vol. 1, Wiley Interscience (Greene 1998)). T cells that have been activated in response to a lung tumor polypeptide, polynucleotide or polypeptide-expressing APC may be CD4⁺ and/or CD8⁺. Lung tumor protein-specific T cells may be expanded using standard techniques. Within preferred embodiments, the T

cells are derived from a patient, a related donor or an unrelated donor, and are administered to the patient following stimulation and expansion.

For therapeutic purposes, CD4⁺ or CD8⁺ T cells that proliferate in response to a lung tumor polypeptide, polynucleotide or APC can be expanded in number either *in vitro* or *in vivo*. Proliferation of such T cells *in vitro* may be accomplished in a variety of ways. For example, the T cells can be re-exposed to a lung tumor polypeptide, or a short peptide corresponding to an immunogenic portion of such a polypeptide, with or without the addition of T cell growth factors, such as interleukin-2, and/or stimulator cells that synthesize a lung tumor polypeptide. Alternatively, one or more T cells that proliferate in the presence of a lung tumor protein can be expanded in number by cloning. Methods for cloning cells are well known in the art, and include limiting dilution.

PHARMACEUTICAL COMPOSITIONS

In additional embodiments, the present invention concerns formulation of one or more of the polynucleotide, polypeptide, T-cell and/or antibody compositions disclosed herein in pharmaceutically-acceptable solutions for administration to a cell or an animal, either alone, or in combination with one or more other modalities of therapy.

It will also be understood that, if desired, the nucleic acid segment, RNA, DNA or PNA compositions that express a polypeptide as disclosed herein may be administered in combination with other agents as well, such as, *e.g.*, other proteins or polypeptides or various pharmaceutically-active agents. In fact, there is virtually no limit to other components that may also be included, given that the additional agents do not cause a significant adverse effect upon contact with the target cells or host tissues. The compositions may thus be delivered along with various other agents as required in the particular instance. Such compositions may be purified from host cells or other biological sources, or alternatively may be chemically synthesized as described herein. Likewise, such compositions may further comprise substituted or derivatized RNA or DNA compositions.

Formulation of pharmaceutically-acceptable excipients and carrier solutions is well-known to those of skill in the art, as is the development of suitable dosing and treatment regimens for using the particular compositions described herein in a variety of treatment regimens, including *e.g.*, oral, parenteral, intravenous, intranasal, and intramuscular administration and formulation.

1. ORAL DELIVERY

In certain applications, the pharmaceutical compositions disclosed herein may be delivered *via* oral administration to an animal. As such, these compositions may be formulated with an inert diluent or with an assimilable edible carrier, or they may be enclosed in hard- or soft-shell gelatin capsule, or they may be compressed into tablets, or they may be incorporated directly with the food of the diet.

The active compounds may even be incorporated with excipients and used in the form of ingestible tablets, buccal tables, troches, capsules, elixirs, suspensions, syrups, wafers, and the like (Mathiowitz *et al.*, 1997; Hwang *et al.*, 1998; U. S. Patent 5,641,515; U. S. Patent 5,580,579 and U. S. Patent 5,792,451, each specifically incorporated herein by reference in its entirety). The tablets, troches, pills, capsules and the like may also contain the following: a binder, as gum tragacanth, acacia, cornstarch, or gelatin; excipients, such as dicalcium phosphate; a disintegrating agent, such as corn starch, potato starch, alginic acid and the like; a lubricant, such as magnesium stearate; and a sweetening agent, such as sucrose, lactose or saccharin may be added or a flavoring agent, such as peppermint, oil of wintergreen, or cherry flavoring. When the dosage unit form is a capsule, it may contain, in addition to materials of the above type, a liquid carrier. Various other materials may be present as coatings or to otherwise modify the physical form of the dosage unit. For instance, tablets, pills, or capsules may be coated with shellac, sugar, or both. A syrup or elixir may contain the active compound sucrose as a sweetening agent methyl and propylparabens as preservatives, a dye and flavoring, such as cherry or orange flavor. Of course, any material used in preparing any dosage unit form should be pharmaceutically pure and substantially non-toxic in the amounts employed. In addition,

the active compounds may be incorporated into sustained-release preparation and formulations.

Typically, these formulations may contain at least about 0.1% of the active compound or more, although the percentage of the active ingredient(s) may, of course, be varied and may conveniently be between about 1 or 2% and about 60% or 70% or more of the weight or volume of the total formulation. Naturally, the amount of active compound(s) in each therapeutically useful composition may be prepared in such a way that a suitable dosage will be obtained in any given unit dose of the compound. Factors such as solubility, bioavailability, biological half-life, route of administration, product shelf life, as well as other pharmacological considerations will be contemplated by one skilled in the art of preparing such pharmaceutical formulations, and as such, a variety of dosages and treatment regimens may be desirable.

For oral administration the compositions of the present invention may alternatively be incorporated with one or more excipients in the form of a mouthwash, dentifrice, buccal tablet, oral spray, or sublingual orally-administered formulation. For example, a mouthwash may be prepared incorporating the active ingredient in the required amount in an appropriate solvent, such as a sodium borate solution (Dobell's Solution). Alternatively, the active ingredient may be incorporated into an oral solution such as one containing sodium borate, glycerin and potassium bicarbonate, or dispersed in a dentifrice, or added in a therapeutically-effective amount to a composition that may include water, binders, abrasives, flavoring agents, foaming agents, and humectants. Alternatively the compositions may be fashioned into a tablet or solution form that may be placed under the tongue or otherwise dissolved in the mouth.

2. INJECTABLE DELIVERY

In certain circumstances it will be desirable to deliver the pharmaceutical compositions disclosed herein parenterally, intravenously, intramuscularly, or even intraperitoneally as described in U. S. Patent 5,543,158; U. S. Patent 5,641,515 and U. S. Patent 5,399,363 (each specifically incorporated herein by reference in its entirety).

Solutions of the active compounds as free base or pharmacologically acceptable salts may be prepared in water suitably mixed with a surfactant, such as hydroxypropylcellulose. Dispersions may also be prepared in glycerol, liquid polyethylene glycols, and mixtures thereof and in oils. Under ordinary conditions of storage and use, these preparations
 5 contain a preservative to prevent the growth of microorganisms.

The pharmaceutical forms suitable for injectable use include sterile aqueous solutions or dispersions and sterile powders for the extemporaneous preparation of sterile injectable solutions or dispersions (U. S. Patent 5,466,468, specifically incorporated herein by reference in its entirety). In all cases the form must be sterile and must be fluid to the
 10 extent that easy syringability exists. It must be stable under the conditions of manufacture and storage and must be preserved against the contaminating action of microorganisms, such as bacteria and fungi. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (*e.g.*, glycerol, propylene glycol, and liquid polyethylene glycol, and the like), suitable mixtures thereof, and/or vegetable oils. Proper
 15 fluidity may be maintained, for example, by the use of a coating, such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. The prevention of the action of microorganisms can be facilitated by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, sorbic acid, thimerosal, and the like. In many cases, it will be preferable to include isotonic
 20 agents, for example, sugars or sodium chloride. Prolonged absorption of the injectable compositions can be brought about by the use in the compositions of agents delaying absorption, for example, aluminum monostearate and gelatin.

For parenteral administration in an aqueous solution, for example, the solution should be suitably buffered if necessary and the liquid diluent first rendered
 25 isotonic with sufficient saline or glucose. These particular aqueous solutions are especially suitable for intravenous, intramuscular, subcutaneous and intraperitoneal administration. In this connection, a sterile aqueous medium that can be employed will be known to those of skill in the art in light of the present disclosure. For example, one dosage may be dissolved in 1 ml of isotonic NaCl solution and either added to 1000 ml of hypodermoclysis fluid or

injected at the proposed site of infusion, (see for example, "Remington's Pharmaceutical Sciences" 15th Edition, pages 1035-1038 and 1570-1580). Some variation in dosage will necessarily occur depending on the condition of the subject being treated. The person responsible for administration will, in any event, determine the appropriate dose for the individual subject. Moreover, for human administration, preparations should meet sterility, pyrogenicity, and the general safety and purity standards as required by FDA Office of Biologics standards.

Sterile injectable solutions are prepared by incorporating the active compounds in the required amount in the appropriate solvent with various of the other ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the various sterilized active ingredients into a sterile vehicle which contains the basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum-drying and freeze-drying techniques which yield a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof.

The compositions disclosed herein may be formulated in a neutral or salt form. Pharmaceutically-acceptable salts, include the acid addition salts (formed with the free amino groups of the protein) and which are formed with inorganic acids such as, for example, hydrochloric or phosphoric acids, or such organic acids as acetic, oxalic, tartaric, mandelic, and the like. Salts formed with the free carboxyl groups can also be derived from inorganic bases such as, for example, sodium, potassium, ammonium, calcium, or ferric hydroxides, and such organic bases as isopropylamine, trimethylamine, histidine, procaine and the like. Upon formulation, solutions will be administered in a manner compatible with the dosage formulation and in such amount as is therapeutically effective. The formulations are easily administered in a variety of dosage forms such as injectable solutions, drug-release capsules, and the like.

As used herein, "carrier" includes any and all solvents, dispersion media, vehicles, coatings, diluents, antibacterial and antifungal agents, isotonic and absorption

delaying agents, buffers, carrier solutions, suspensions, colloids, and the like. The use of such media and agents for pharmaceutical active substances is well known in the art. Except insofar as any conventional media or agent is incompatible with the active ingredient, its use in the therapeutic compositions is contemplated. Supplementary active
 5 ingredients can also be incorporated into the compositions.

The phrase "pharmaceutically-acceptable" refers to molecular entities and compositions that do not produce an allergic or similar untoward reaction when administered to a human. The preparation of an aqueous composition that contains a protein as an active ingredient is well understood in the art. Typically, such compositions
 10 are prepared as injectables, either as liquid solutions or suspensions; solid forms suitable for solution in, or suspension in, liquid prior to injection can also be prepared. The preparation can also be emulsified.

3. NASAL DELIVERY

In certain embodiments, the pharmaceutical compositions may be delivered
 15 by intranasal sprays, inhalation, and/or other aerosol delivery vehicles. Methods for delivering genes, nucleic acids, and peptide compositions directly to the lungs *via* nasal aerosol sprays has been described *e.g.*, in U. S. Patent 5,756,353 and U. S. Patent 5,804,212 (each specifically incorporated herein by reference in its entirety). Likewise, the delivery of drugs using intranasal microparticle resins (Takenaga *et al.*, 1998) and lysophosphatidyl-
 20 glycerol compounds (U. S. Patent 5,725,871, specifically incorporated herein by reference in its entirety) are also well-known in the pharmaceutical arts. Likewise, transmucosal drug delivery in the form of a polytetrafluoroethylene support matrix is described in U. S. Patent 5,780,045 (specifically incorporated herein by reference in its entirety).

4. LIPOSOME-, NANOCAPSULE-, AND MICROPARTICLE-MEDIATED DELIVERY

25 In certain embodiments, the inventors contemplate the use of liposomes, nanocapsules, microparticles, microspheres, lipid particles, vesicles, and the like, for the introduction of the compositions of the present invention into suitable host cells. In

particular, the compositions of the present invention may be formulated for delivery either encapsulated in a lipid particle, a liposome, a vesicle, a nanosphere, or a nanoparticle or the like.

Such formulations may be preferred for the introduction of
 5 pharmaceutically-acceptable formulations of the nucleic acids or constructs disclosed herein. The formation and use of liposomes is generally known to those of skill in the art (see for example, Couvreur *et al.*, 1977; Couvreur, 1988; Lasic, 1998; which describes the use of liposomes and nanocapsules in the targeted antibiotic therapy for intracellular bacterial infections and diseases). Recently, liposomes were developed with improved
 10 serum stability and circulation half-times (Gabizon and Papahadjopoulos, 1988; Allen and Choun, 1987; U. S. Patent 5,741,516, specifically incorporated herein by reference in its entirety). Further, various methods of liposome and liposome like preparations as potential drug carriers have been reviewed (Takakura, 1998; Chandran *et al.*, 1997; Margalit, 1995; U. S. Patent 5,567,434; U. S. Patent 5,552,157; U. S. Patent 5,565,213; U. S. Patent
 15 5,738,868 and U. S. Patent 5,795,587, each specifically incorporated herein by reference in its entirety).

Liposomes have been used successfully with a number of cell types that are normally resistant to transfection by other procedures including T cell suspensions, primary hepatocyte cultures and PC 12 cells (Renneisen *et al.*, 1990; Muller *et al.*, 1990). In
 20 addition, liposomes are free of the DNA length constraints that are typical of viral-based delivery systems. Liposomes have been used effectively to introduce genes, drugs (Heath and Martin, 1986; Heath *et al.*, 1986; Balazsovits *et al.*, 1989; Fresta and Puglisi, 1996), radiotherapeutic agents (Pikul *et al.*, 1987), enzymes (Imaizumi *et al.*, 1990a; Imaizumi
 25 *et al.*, 1990b), viruses (Faller and Baltimore, 1984), transcription factors and allosteric effectors (Nicolau and Gersonde, 1979) into a variety of cultured cell lines and animals. In addition, several successful clinical trails examining the effectiveness of liposome-mediated drug delivery have been completed (Lopez-Berestein *et al.*, 1985a; 1985b; Coune, 1988; Sculier *et al.*, 1988). Furthermore, several studies suggest that the use of

liposomes is not associated with autoimmune responses, toxicity or gonadal localization after systemic delivery (Mori and Fukatsu, 1992).

Liposomes are formed from phospholipids that are dispersed in an aqueous medium and spontaneously form multilamellar concentric bilayer vesicles (also termed
 5 multilamellar vesicles (MLVs). MLVs generally have diameters of from 25 nm to 4 μ m. Sonication of MLVs results in the formation of small unilamellar vesicles (SUVs) with diameters in the range of 200 to 500 Å, containing an aqueous solution in the core.

Liposomes bear resemblance to cellular membranes and are contemplated for use in connection with the present invention as carriers for the peptide compositions.
 10 They are widely suitable as both water- and lipid-soluble substances can be entrapped, *i.e.* in the aqueous spaces and within the bilayer itself, respectively. It is possible that the drug-bearing liposomes may even be employed for site-specific delivery of active agents by selectively modifying the liposomal formulation.

In addition to the teachings of Couvreur *et al.* (1977; 1988), the following
 15 information may be utilized in generating liposomal formulations. Phospholipids can form a variety of structures other than liposomes when dispersed in water, depending on the molar ratio of lipid to water. At low ratios the liposome is the preferred structure. The physical characteristics of liposomes depend on pH, ionic strength and the presence of divalent cations. Liposomes can show low permeability to ionic and polar substances, but
 20 at elevated temperatures undergo a phase transition which markedly alters their permeability. The phase transition involves a change from a closely packed, ordered structure, known as the gel state, to a loosely packed, less-ordered structure, known as the fluid state. This occurs at a characteristic phase-transition temperature and results in an increase in permeability to ions, sugars and drugs.

25 In addition to temperature, exposure to proteins can alter the permeability of liposomes. Certain soluble proteins, such as cytochrome c, bind, deform and penetrate the bilayer, thereby causing changes in permeability. Cholesterol inhibits this penetration of proteins, apparently by packing the phospholipids more tightly. It is contemplated that the

most useful liposome formations for antibiotic and inhibitor delivery will contain cholesterol.

The ability to trap solutes varies between different types of liposomes. For example, MLVs are moderately efficient at trapping solutes, but SUVs are extremely inefficient. SUVs offer the advantage of homogeneity and reproducibility in size distribution, however, and a compromise between size and trapping efficiency is offered by large unilamellar vesicles (LUVs). These are prepared by ether evaporation and are three to four times more efficient at solute entrapment than MLVs.

In addition to liposome characteristics, an important determinant in entrapping compounds is the physicochemical properties of the compound itself. Polar compounds are trapped in the aqueous spaces and nonpolar compounds bind to the lipid bilayer of the vesicle. Polar compounds are released through permeation or when the bilayer is broken, but nonpolar compounds remain affiliated with the bilayer unless it is disrupted by temperature or exposure to lipoproteins. Both types show maximum efflux rates at the phase transition temperature.

Liposomes interact with cells *via* four different mechanisms: endocytosis by phagocytic cells of the reticuloendothelial system such as macrophages and neutrophils; adsorption to the cell surface, either by nonspecific weak hydrophobic or electrostatic forces, or by specific interactions with cell-surface components; fusion with the plasma cell membrane by insertion of the lipid bilayer of the liposome into the plasma membrane, with simultaneous release of liposomal contents into the cytoplasm; and by transfer of liposomal lipids to cellular or subcellular membranes, or vice versa, without any association of the liposome contents. It often is difficult to determine which mechanism is operative and more than one may operate at the same time.

The fate and disposition of intravenously injected liposomes depend on their physical properties, such as size, fluidity, and surface charge. They may persist in tissues for h or days, depending on their composition, and half lives in the blood range from min to several h. Larger liposomes, such as MLVs and LUVs, are taken up rapidly by phagocytic cells of the reticuloendothelial system, but physiology of the circulatory system restrains

the exit of such large species at most sites. They can exit only in places where large openings or pores exist in the capillary endothelium, such as the sinusoids of the liver or spleen. Thus, these organs are the predominate site of uptake. On the other hand, SUVs show a broader tissue distribution but still are sequestered highly in the liver and spleen. In
 5 general, this *in vivo* behavior limits the potential targeting of liposomes to only those organs and tissues accessible to their large size. These include the blood, liver, spleen, bone marrow, and lymphoid organs.

Targeting is generally not a limitation in terms of the present invention. However, should specific targeting be desired, methods are available for this to be
 10 accomplished. Antibodies may be used to bind to the liposome surface and to direct the antibody and its drug contents to specific antigenic receptors located on a particular cell-type surface. Carbohydrate determinants (glycoprotein or glycolipid cell-surface components that play a role in cell-cell recognition, interaction and adhesion) may also be used as recognition sites as they have potential in directing liposomes to particular cell
 15 types. Mostly, it is contemplated that intravenous injection of liposomal preparations would be used, but other routes of administration are also conceivable.

Alternatively, the invention provides for pharmaceutically-acceptable nanocapsule formulations of the compositions of the present invention. Nanocapsules can generally entrap compounds in a stable and reproducible way (Henry-Michelland *et al.*,
 20 1987; Quintanar-Guerrero *et al.*, 1998; Douglas *et al.*, 1987). To avoid side effects due to intracellular polymeric overloading, such ultrafine particles (sized around 0.1 μm) should be designed using polymers able to be degraded *in vivo*. Biodegradable polyalkyl-cyanoacrylate nanoparticles that meet these requirements are contemplated for use in the present invention. Such particles may be easily made, as described (Couvreur *et al.*,
 25 1980; 1988; zur Muhlen *et al.*, 1998; Zambaux *et al.* 1998; Pinto-Alphandry *et al.*, 1995 and U. S. Patent 5,145,684, specifically incorporated herein by reference in its entirety).

IMMUNOGENIC COMPOSITIONS

In certain preferred embodiments of the present invention, immunogenic compositions, or vaccines, are provided. The immunogenic compositions will generally comprise one or more pharmaceutical compositions, such as those discussed above, in combination with an immunostimulant. An immunostimulant may be any substance that enhances or potentiates an immune response (antibody and/or cell-mediated) to an exogenous antigen. Examples of immunostimulants include adjuvants, biodegradable microspheres (*e.g.*, polylactic galactide) and liposomes (into which the compound is incorporated; *see e.g.*, Fullerton, U.S. Patent No. 4,235,877). Vaccine preparation is generally described in, for example, M.F. Powell and M.J. Newman, eds., "Vaccine Design (the subunit and adjuvant approach)," Plenum Press (NY, 1995). Pharmaceutical compositions and immunogenic compositions, or vaccines, within the scope of the present invention may also contain other compounds, which may be biologically active or inactive. For example, one or more immunogenic portions of other tumor antigens may be present, either incorporated into a fusion polypeptide or as a separate compound, within the composition.

Illustrative immunogenic compositions may contain DNA encoding one or more of the polypeptides as described above, such that the polypeptide is generated *in situ*. As noted above, the DNA may be present within any of a variety of delivery systems known to those of ordinary skill in the art, including nucleic acid expression systems, bacteria and viral expression systems. Numerous gene delivery techniques are well known in the art, such as those described by Rolland, *Crit. Rev. Therap. Drug Carrier Systems* 15:143-198, 1998, and references cited therein. Appropriate nucleic acid expression systems contain the necessary DNA sequences for expression in the patient (such as a suitable promoter and terminating signal). Bacterial delivery systems involve the administration of a bacterium (such as *Bacillus-Calmette-Guerrin*) that expresses an immunogenic portion of the polypeptide on its cell surface or secretes such an epitope. In a preferred embodiment, the DNA may be introduced using a viral expression system (*e.g.*, vaccinia or other pox virus, retrovirus, or adenovirus), which may involve the use of a non-

pathogenic (defective), replication competent virus. Suitable systems are disclosed, for example, in Fisher-Hoch *et al.*, *Proc. Natl. Acad. Sci. USA* 86:317-321, 1989; Flexner *et al.*, *Ann. N.Y. Acad. Sci.* 569:86-103, 1989; Flexner *et al.*, *Vaccine* 8:17-21, 1990; U.S. Patent Nos. 4,603,112, 4,769,330, and 5,017,487; WO 89/01973; U.S. Patent
 5 No. 4,777,127; GB 2,200,651; EP 0,345,242; WO 91/02805; Berkner, *Biotechniques* 6:616-627, 1988; Rosenfeld *et al.*, *Science* 252:431-434, 1991; Kolls *et al.*, *Proc. Natl. Acad. Sci. USA* 91:215-219, 1994; Kass-Eisler *et al.*, *Proc. Natl. Acad. Sci. USA* 90:11498-11502, 1993; Guzman *et al.*, *Circulation* 88:2838-2848, 1993; and Guzman *et al.*, *Cir. Res.* 73:1202-1207, 1993. Techniques for incorporating DNA into such expression
 10 systems are well known to those of ordinary skill in the art. The DNA may also be "naked," as described, for example, in Ulmer *et al.*, *Science* 259:1745-1749, 1993 and reviewed by Cohen, *Science* 259:1691-1692, 1993. The uptake of naked DNA may be increased by coating the DNA onto biodegradable beads, which are efficiently transported into the cells. It will be apparent that an immunogenic composition may comprise both a
 15 polynucleotide and a polypeptide component. Such immunogenic compositions may provide for an enhanced immune response.

It will be apparent that an immunogenic composition may contain pharmaceutically acceptable salts of the polynucleotides and polypeptides provided herein. Such salts may be prepared from pharmaceutically acceptable non-toxic bases, including
 20 organic bases (*e.g.*, salts of primary, secondary and tertiary amines and basic amino acids) and inorganic bases (*e.g.*, sodium, potassium, lithium, ammonium, calcium and magnesium salts).

While any suitable carrier known to those of ordinary skill in the art may be employed in the immunogenic compositions of this invention, the type of carrier will vary
 25 depending on the mode of administration. Compositions of the present invention may be formulated for any appropriate manner of administration, including for example, topical, oral, nasal, intravenous, intracranial, intraperitoneal, subcutaneous or intramuscular administration. For parenteral administration, such as subcutaneous injection, the carrier preferably comprises water, saline, alcohol, a fat, a wax or a buffer. For oral

administration, any of the above carriers or a solid carrier, such as mannitol, lactose, starch, magnesium stearate, sodium saccharine, talcum, cellulose, glucose, sucrose, and magnesium carbonate, may be employed. Biodegradable microspheres (*e.g.*, polylactate polyglycolate) may also be employed as carriers for the pharmaceutical compositions of this invention. Suitable biodegradable microspheres are disclosed, for example, in U.S. Patent Nos. 4,897,268; 5,075,109; 5,928,647; 5,811,128; 5,820,883; 5,853,763; 5,814,344 and 5,942,252. One may also employ a carrier comprising the particulate-protein complexes described in U.S. Patent No. 5,928,647, which are capable of inducing a class I-restricted cytotoxic T lymphocyte responses in a host.

Such compositions may also comprise buffers (*e.g.*, neutral buffered saline or phosphate buffered saline), carbohydrates (*e.g.*, glucose, mannose, sucrose or dextrans), mannitol, proteins, polypeptides or amino acids such as glycine, antioxidants, bacteriostats, chelating agents such as EDTA or glutathione, adjuvants (*e.g.*, aluminum hydroxide), solutes that render the formulation isotonic, hypotonic or weakly hypertonic with the blood of a recipient, suspending agents, thickening agents and/or preservatives. Alternatively, compositions of the present invention may be formulated as a lyophilizate. Compounds may also be encapsulated within liposomes using well known technology.

Any of a variety of immunostimulants may be employed in the immunogenic compositions of this invention. For example, an adjuvant may be included. Most adjuvants contain a substance designed to protect the antigen from rapid catabolism, such as aluminum hydroxide or mineral oil, and a stimulator of immune responses, such as lipid A, *Bordetella pertussis* or *Mycobacterium tuberculosis* derived proteins. Suitable adjuvants are commercially available as, for example, Freund's Incomplete Adjuvant and Complete Adjuvant (Difco Laboratories, Detroit, MI); Merck Adjuvant 65 (Merck and Company, Inc., Rahway, NJ); AS-2 (SmithKline Beecham, Philadelphia, PA); aluminum salts such as aluminum hydroxide gel (alum) or aluminum phosphate; salts of calcium, iron or zinc; an insoluble suspension of acylated tyrosine; acylated sugars; cationically or anionically derivatized polysaccharides; polyphosphazenes; biodegradable microspheres;

monophosphoryl lipid A and quil A. Cytokines, such as GM-CSF or interleukin-2, -7, or -12, may also be used as adjuvants.

Within the immunogenic compositions provided herein, the adjuvant composition is preferably designed to induce an immune response predominantly of the Th1 type. High levels of Th1-type cytokines (*e.g.*, IFN- γ , TNF α , IL-2 and IL-12) tend to favor the induction of cell mediated immune responses to an administered antigen. In contrast, high levels of Th2-type cytokines (*e.g.*, IL-4, IL-5, IL-6 and IL-10) tend to favor the induction of humoral immune responses. Following application of an immunogenic composition as provided herein, a patient will support an immune response that includes Th1- and Th2-type responses. Within a preferred embodiment, in which a response is predominantly Th1-type, the level of Th1-type cytokines will increase to a greater extent than the level of Th2-type cytokines. The levels of these cytokines may be readily assessed using standard assays. For a review of the families of cytokines, see Mosmann and Coffman, *Ann. Rev. Immunol.* 7:145-173, 1989.

Preferred adjuvants for use in eliciting a predominantly Th1-type response include, for example, a combination of monophosphoryl lipid A, preferably 3-de-O-acylated monophosphoryl lipid A (3D-MPL), together with an aluminum salt. MPL adjuvants are available from Corixa Corporation (Seattle, WA; *see* US Patent Nos. 4,436,727; 4,877,611; 4,866,034 and 4,912,094). CpG-containing oligonucleotides (in which the CpG dinucleotide is unmethylated) also induce a predominantly Th1 response. Such oligonucleotides are well known and are described, for example, in WO 96/02555, WO 99/33488 and U.S. Patent Nos. 6,008,200 and 5,856,462. Immunostimulatory DNA sequences are also described, for example, by Sato *et al.*, *Science* 273:352, 1996. Another preferred adjuvant is a saponin, preferably QS21 (Aquila Biopharmaceuticals Inc., Framingham, MA), which may be used alone or in combination with other adjuvants. For example, an enhanced system involves the combination of a monophosphoryl lipid A and saponin derivative, such as the combination of QS21 and 3D-MPL as described in WO 94/00153, or a less reactogenic composition where the QS21 is quenched with cholesterol, as described in WO 96/33739. Other preferred formulations comprise an oil-in-water

emulsion and tocopherol. A particularly potent adjuvant formulation involving QS21, 3D-MPL and tocopherol in an oil-in-water emulsion is described in WO 95/17210.

Other preferred adjuvants include Montanide ISA 720 (Seppic, France), SAF (Chiron, California, United States), ISCOMS (CSL), MF-59 (Chiron), the SBAS series of adjuvants (*e.g.*, SBAS-2 or SBAS-4, available from SmithKline Beecham, Rixensart, Belgium), Detox (Corixa, Hamilton, MT), RC-529 (Corixa, Hamilton, MT) and other aminoalkyl glucosaminide 4-phosphates (AGPs), such as those described in pending U.S. Patent Application Serial Nos. 08/853,826 and 09/074,720, the disclosures of which are incorporated herein by reference in their entireties.

Any immunogenic composition provided herein may be prepared using well known methods that result in a combination of antigen, immune response enhancer and a suitable carrier or excipient. The compositions described herein may be administered as part of a sustained release formulation (*i.e.*, a formulation such as a capsule, sponge or gel (composed of polysaccharides, for example) that effects a slow release of compound following administration). Such formulations may generally be prepared using well known technology (*see, e.g.*, Coombes *et al.*, *Vaccine* 14:1429-1438, 1996) and administered by, for example, oral, rectal or subcutaneous implantation, or by implantation at the desired target site. Sustained-release formulations may contain a polypeptide, polynucleotide or antibody dispersed in a carrier matrix and/or contained within a reservoir surrounded by a rate controlling membrane.

Carriers for use within such formulations are biocompatible, and may also be biodegradable; preferably the formulation provides a relatively constant level of active component release. Such carriers include microparticles of poly(lactide-co-glycolide), polyacrylate, latex, starch, cellulose, dextran and the like. Other delayed-release carriers include supramolecular biovectors, which comprise a non-liquid hydrophilic core (*e.g.*, a cross-linked polysaccharide or oligosaccharide) and, optionally, an external layer comprising an amphiphilic compound, such as a phospholipid (*see e.g.*, U.S. Patent No. 5,151,254 and PCT applications WO 94/20078, WO/94/23701 and WO 96/06638). The amount of active compound contained within a sustained release formulation depends upon

the site of implantation, the rate and expected duration of release and the nature of the condition to be treated or prevented.

Any of a variety of delivery vehicles may be employed within pharmaceutical compositions and immunogenic compositions to facilitate production of an antigen-specific immune response that targets tumor cells. Delivery vehicles include antigen presenting cells (APCs), such as dendritic cells, macrophages, B cells, monocytes and other cells that may be engineered to be efficient APCs. Such cells may, but need not, be genetically modified to increase the capacity for presenting the antigen, to improve activation and/or maintenance of the T cell response, to have anti-tumor effects *per se* and/or to be immunologically compatible with the receiver (*i.e.*, matched HLA haplotype). APCs may generally be isolated from any of a variety of biological fluids and organs, including tumor and peritumoral tissues, and may be autologous, allogeneic, syngeneic or xenogeneic cells.

Certain preferred embodiments of the present invention use dendritic cells or progenitors thereof as antigen-presenting cells. Dendritic cells are highly potent APCs (Banchereau and Steinman, *Nature* 392:245-251, 1998) and have been shown to be effective as a physiological adjuvant for eliciting prophylactic or therapeutic antitumor immunity (*see* Timmerman and Levy, *Ann. Rev. Med.* 50:507-529, 1999). In general, dendritic cells may be identified based on their typical shape (stellate *in situ*, with marked cytoplasmic processes (dendrites) visible *in vitro*), their ability to take up, process and present antigens with high efficiency and their ability to activate naïve T cell responses. Dendritic cells may, of course, be engineered to express specific cell-surface receptors or ligands that are not commonly found on dendritic cells *in vivo* or *ex vivo*, and such modified dendritic cells are contemplated by the present invention. As an alternative to dendritic cells, secreted vesicles antigen-loaded dendritic cells (called exosomes) may be used within a vaccine, or immunogenic composition (*see* Zitvogel *et al.*, *Nature Med.* 4:594-600, 1998).

Dendritic cells and progenitors may be obtained from peripheral blood, bone marrow, tumor-infiltrating cells, peritumoral tissues-infiltrating cells, lymph nodes, spleen,

skin, umbilical cord blood or any other suitable tissue or fluid. For example, dendritic cells may be differentiated *ex vivo* by adding a combination of cytokines such as GM-CSF, IL-4, IL-13 and/or TNF α to cultures of monocytes harvested from peripheral blood. Alternatively, CD34 positive cells harvested from peripheral blood, umbilical cord blood or
 5 bone marrow may be differentiated into dendritic cells by adding to the culture medium combinations of GM-CSF, IL-3, TNF α , CD40 ligand, LPS, flt3 ligand and/or other compound(s) that induce differentiation, maturation and proliferation of dendritic cells.

Dendritic cells are conveniently categorized as "immature" and "mature" cells, which allows a simple way to discriminate between two well characterized
 10 phenotypes. However, this nomenclature should not be construed to exclude all possible intermediate stages of differentiation. Immature dendritic cells are characterized as APC with a high capacity for antigen uptake and processing, which correlates with the high expression of Fc γ receptor and mannose receptor. The mature phenotype is typically characterized by a lower expression of these markers, but a high expression of cell surface
 15 molecules responsible for T cell activation such as class I and class II MHC, adhesion molecules (*e.g.*, CD54 and CD11) and costimulatory molecules (*e.g.*, CD40, CD80, CD86 and 4-1BB).

APCs may generally be transfected with a polynucleotide encoding a lung tumor protein (or portion or other variant thereof) such that the lung tumor polypeptide, or
 20 an immunogenic portion thereof, is expressed on the cell surface. Such transfection may take place *ex vivo*, and a composition comprising such transfected cells may then be used for therapeutic purposes, as described herein. Alternatively, a gene delivery vehicle that targets a dendritic or other antigen presenting cell may be administered to a patient, resulting in transfection that occurs *in vivo*. *In vivo* and *ex vivo* transfection of dendritic
 25 cells, for example, may generally be performed using any methods known in the art, such as those described in WO 97/24447, or the gene gun approach described by Mahvi *et al.*, *Immunology and cell Biology* 75:456-460, 1997. Antigen loading of dendritic cells may be achieved by incubating dendritic cells or progenitor cells with the lung tumor polypeptide, DNA (naked or within a plasmid vector) or RNA; or with antigen-expressing recombinant

bacterium or viruses (*e.g.*, vaccinia, fowlpox, adenovirus or lentivirus vectors). Prior to loading, the polypeptide may be covalently conjugated to an immunological partner that provides T cell help (*e.g.*, a carrier molecule). Alternatively, a dendritic cell may be pulsed with a non-conjugated immunological partner, separately or in the presence of the polypeptide.

Immunogenic compositions and pharmaceutical compositions may be presented in unit-dose or multi-dose containers, such as sealed ampoules or vials. Such containers are preferably hermetically sealed to preserve sterility of the formulation until use. In general, formulations may be stored as suspensions, solutions or emulsions in oily or aqueous vehicles. Alternatively, an immunogenic or pharmaceutical composition may be stored in a freeze-dried condition requiring only the addition of a sterile liquid carrier immediately prior to use.

CANCER THERAPY

In further aspects of the present invention, the compositions described herein may be used for immunotherapy of cancer, such as lung cancer. Within such methods, compositions are typically administered to a patient. As used herein, a “patient” refers to any warm-blooded animal, preferably a human. A patient may or may not be afflicted with cancer. Accordingly, the above pharmaceutical compositions and immunogenic compositions may be used to prevent the development of a cancer or to treat a patient afflicted with a cancer. A cancer may be diagnosed using criteria generally accepted in the art, including the presence of a malignant tumor. Pharmaceutical compositions and immunogenic compositions may be administered either prior to or following surgical removal of primary tumors and/or treatment such as administration of radiotherapy or conventional chemotherapeutic drugs. Administration may be by any suitable method, including administration by intravenous, intraperitoneal, intramuscular, subcutaneous, intranasal, intradermal, anal, vaginal, topical and oral routes.

Within certain embodiments, immunotherapy may be active immunotherapy, in which treatment relies on the *in vivo* stimulation of the endogenous host

immune system to react against tumors with the administration of immune response-modifying agents (such as polypeptides and polynucleotides as provided herein).

Within other embodiments, immunotherapy may be passive immunotherapy, in which treatment involves the delivery of agents with established tumor-immune reactivity (such as effector cells or antibodies) that can directly or indirectly mediate antitumor effects and does not necessarily depend on an intact host immune system. Examples of effector cells include T cells as discussed above, T lymphocytes (such as CD8⁺ cytotoxic T lymphocytes and CD4⁺ T-helper tumor-infiltrating lymphocytes), killer cells (such as Natural Killer cells and lymphokine-activated killer cells), B cells and antigen-presenting cells (such as dendritic cells and macrophages) expressing a polypeptide provided herein. T cell receptors and antibody receptors specific for the polypeptides recited herein may be cloned, expressed and transferred into other vectors or effector cells for adoptive immunotherapy. The polypeptides provided herein may also be used to generate antibodies or anti-idiotypic antibodies (as described above and in U.S. Patent No. 4,918,164) for passive immunotherapy.

Effector cells may generally be obtained in sufficient quantities for adoptive immunotherapy by growth *in vitro*, as described herein. Culture conditions for expanding single antigen-specific effector cells to several billion in number with retention of antigen recognition *in vivo* are well known in the art. Such *in vitro* culture conditions typically use intermittent stimulation with antigen, often in the presence of cytokines (such as IL-2) and non-dividing feeder cells. As noted above, immunoreactive polypeptides as provided herein may be used to rapidly expand antigen-specific T cell cultures in order to generate a sufficient number of cells for immunotherapy. In particular, antigen-presenting cells, such as dendritic, macrophage, monocyte, fibroblast and/or B cells, may be pulsed with immunoreactive polypeptides or transfected with one or more polynucleotides using standard techniques well known in the art. For example, antigen-presenting cells can be transfected with a polynucleotide having a promoter appropriate for increasing expression in a recombinant virus or other expression system. Cultured effector cells for use in therapy must be able to grow and distribute widely, and to survive long term *in vivo*.

Studies have shown that cultured effector cells can be induced to grow *in vivo* and to survive long term in substantial numbers by repeated stimulation with antigen supplemented with IL-2 (*see, for example, Cheever et al., Immunological Reviews 157:177, 1997*).

5 Alternatively, a vector expressing a polypeptide recited herein may be introduced into antigen presenting cells taken from a patient and clonally propagated *ex vivo* for transplant back into the same patient. Transfected cells may be reintroduced into the patient using any means known in the art, preferably in sterile form by intravenous, intracavitary, intraperitoneal or intratumor administration.

10 Routes and frequency of administration of the therapeutic compositions described herein, as well as dosage, will vary from individual to individual, and may be readily established using standard techniques. In general, the pharmaceutical compositions and immunogenic compositions may be administered by injection (*e.g.*, intracutaneous, intramuscular, intravenous or subcutaneous), intranasally (*e.g.*, by aspiration) or orally.

15 Preferably, between 1 and 10 doses may be administered over a 52 week period. Preferably, 6 doses are administered, at intervals of 1 month, and booster vaccinations may be given periodically thereafter. Alternate protocols may be appropriate for individual patients. A suitable dose is an amount of a compound that, when administered as described above, is capable of promoting an anti-tumor immune response, and is at least 10-50%

20 above the basal (*i.e.*, untreated) level. Such response can be monitored by measuring the anti-tumor antibodies in a patient or by vaccine-dependent generation of cytolytic effector cells capable of killing the patient's tumor cells *in vitro*. Such vaccines, or immunogenic compositions, should also be capable of causing an immune response that leads to an improved clinical outcome (*e.g.*, more frequent remissions, complete or partial or longer

25 disease-free survival) in vaccinated patients as compared to non-vaccinated patients. In general, for compositions comprising one or more polypeptides, the amount of each polypeptide present in a dose ranges from about 25 μ g to 5 mg per kg of host. Suitable dose sizes will vary with the size of the patient, but will typically range from about 0.1 mL to about 5 mL.

In general, an appropriate dosage and treatment regimen provides the active compound(s) in an amount sufficient to provide therapeutic and/or prophylactic benefit. Such a response can be monitored by establishing an improved clinical outcome (*e.g.*, more frequent remissions, complete or partial, or longer disease-free survival) in treated patients as compared to non-treated patients. Increases in preexisting immune responses to a lung tumor protein generally correlate with an improved clinical outcome. Such immune responses may generally be evaluated using standard proliferation, cytotoxicity or cytokine assays, which may be performed using samples obtained from a patient before and after treatment.

10 **CANCER DETECTION AND DIAGNOSIS**

In general, a cancer may be detected in a patient based on the presence of one or more lung tumor proteins and/or polynucleotides encoding such proteins in a biological sample (for example, blood, sera, sputum urine and/or tumor biopsies) obtained from the patient. In other words, such proteins may be used as markers to indicate the presence or absence of a cancer such as lung cancer. In addition, such proteins may be useful for the detection of other cancers. The binding agents provided herein generally permit detection of the level of antigen that binds to the agent in the biological sample. Polynucleotide primers and probes may be used to detect the level of mRNA encoding a tumor protein, which is also indicative of the presence or absence of a cancer. In general, a lung tumor sequence should be present at a level that is at least three fold higher in tumor tissue than in normal tissue

There are a variety of assay formats known to those of ordinary skill in the art for using a binding agent to detect polypeptide markers in a sample. *See, e.g.*, Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. In general, the presence or absence of a cancer in a patient may be determined by (a) contacting a biological sample obtained from a patient with a binding agent; (b) detecting in the sample a level of polypeptide that binds to the binding agent; and (c) comparing the level of polypeptide with a predetermined cut-off value.

In a preferred embodiment, the assay involves the use of binding agent immobilized on a solid support to bind to and remove the polypeptide from the remainder of the sample. The bound polypeptide may then be detected using a detection reagent that contains a reporter group and specifically binds to the binding agent/polypeptide complex.

5 Such detection reagents may comprise, for example, a binding agent that specifically binds to the polypeptide or an antibody or other agent that specifically binds to the binding agent, such as an anti-immunoglobulin, protein G, protein A or a lectin. Alternatively, a competitive assay may be utilized, in which a polypeptide is labeled with a reporter group and allowed to bind to the immobilized binding agent after incubation of the binding agent

10 with the sample. The extent to which components of the sample inhibit the binding of the labeled polypeptide to the binding agent is indicative of the reactivity of the sample with the immobilized binding agent. Suitable polypeptides for use within such assays include full length lung tumor proteins and portions thereof to which the binding agent binds, as described above.

15 The solid support may be any material known to those of ordinary skill in the art to which the tumor protein may be attached. For example, the solid support may be a test well in a microtiter plate or a nitrocellulose or other suitable membrane. Alternatively, the support may be a bead or disc, such as glass, fiberglass, latex or a plastic material such as polystyrene or polyvinylchloride. The support may also be a magnetic

20 particle or a fiber optic sensor, such as those disclosed, for example, in U.S. Patent No. 5,359,681. The binding agent may be immobilized on the solid support using a variety of techniques known to those of skill in the art, which are amply described in the patent and scientific literature. In the context of the present invention, the term "immobilization" refers to both noncovalent association, such as adsorption, and covalent attachment (which

25 may be a direct linkage between the agent and functional groups on the support or may be a linkage by way of a cross-linking agent). Immobilization by adsorption to a well in a microtiter plate or to a membrane is preferred. In such cases, adsorption may be achieved by contacting the binding agent, in a suitable buffer, with the solid support for a suitable amount of time. The contact time varies with temperature, but is typically between about 1

hour and about 1 day. In general, contacting a well of a plastic microtiter plate (such as polystyrene or polyvinylchloride) with an amount of binding agent ranging from about 10 ng to about 10 μ g, and preferably about 100 ng to about 1 μ g, is sufficient to immobilize an adequate amount of binding agent.

5 Covalent attachment of binding agent to a solid support may generally be achieved by first reacting the support with a bifunctional reagent that will react with both the support and a functional group, such as a hydroxyl or amino group, on the binding agent. For example, the binding agent may be covalently attached to supports having an appropriate polymer coating using benzoquinone or by condensation of an aldehyde group
10 on the support with an amine and an active hydrogen on the binding partner (*see, e.g.*, Pierce Immunotechnology Catalog and Handbook, 1991, at A12-A13).

In certain embodiments, the assay is a two-antibody sandwich assay. This assay may be performed by first contacting an antibody that has been immobilized on a solid support, commonly the well of a microtiter plate, with the sample, such that
15 polypeptides within the sample are allowed to bind to the immobilized antibody. Unbound sample is then removed from the immobilized polypeptide-antibody complexes and a detection reagent (preferably a second antibody capable of binding to a different site on the polypeptide) containing a reporter group is added. The amount of detection reagent that remains bound to the solid support is then determined using a method appropriate for the
20 specific reporter group.

More specifically, once the antibody is immobilized on the support as described above, the remaining protein binding sites on the support are typically blocked. Any suitable blocking agent known to those of ordinary skill in the art, such as bovine serum albumin or Tween 20TM (Sigma Chemical Co., St. Louis, MO). The immobilized
25 antibody is then incubated with the sample, and polypeptide is allowed to bind to the antibody. The sample may be diluted with a suitable diluent, such as phosphate-buffered saline (PBS) prior to incubation. In general, an appropriate contact time (*i.e.*, incubation time) is a period of time that is sufficient to detect the presence of polypeptide within a sample obtained from an individual with lung cancer. Preferably, the contact time is

sufficient to achieve a level of binding that is at least about 95% of that achieved at equilibrium between bound and unbound polypeptide. Those of ordinary skill in the art will recognize that the time necessary to achieve equilibrium may be readily determined by assaying the level of binding that occurs over a period of time. At room temperature, an incubation time of about 30 minutes is generally sufficient.

Unbound sample may then be removed by washing the solid support with an appropriate buffer, such as PBS containing 0.1% Tween 20™. The second antibody, which contains a reporter group, may then be added to the solid support. Preferred reporter groups include those groups recited above.

The detection reagent is then incubated with the immobilized antibody-polypeptide complex for an amount of time sufficient to detect the bound polypeptide. An appropriate amount of time may generally be determined by assaying the level of binding that occurs over a period of time. Unbound detection reagent is then removed and bound detection reagent is detected using the reporter group. The method employed for detecting the reporter group depends upon the nature of the reporter group. For radioactive groups, scintillation counting or autoradiographic methods are generally appropriate. Spectroscopic methods may be used to detect dyes, luminescent groups and fluorescent groups. Biotin may be detected using avidin, coupled to a different reporter group (commonly a radioactive or fluorescent group or an enzyme). Enzyme reporter groups may generally be detected by the addition of substrate (generally for a specific period of time), followed by spectroscopic or other analysis of the reaction products.

To determine the presence or absence of a cancer, such as lung cancer, the signal detected from the reporter group that remains bound to the solid support is generally compared to a signal that corresponds to a predetermined cut-off value. In one preferred embodiment, the cut-off value for the detection of a cancer is the average mean signal obtained when the immobilized antibody is incubated with samples from patients without the cancer. In general, a sample generating a signal that is three standard deviations above the predetermined cut-off value is considered positive for the cancer. In an alternate preferred embodiment, the cut-off value is determined using a Receiver Operator Curve,

according to the method of Sackett *et al.*, *Clinical Epidemiology: A Basic Science for Clinical Medicine*, Little Brown and Co., 1985, p. 106-7. Briefly, in this embodiment, the cut-off value may be determined from a plot of pairs of true positive rates (*i.e.*, sensitivity) and false positive rates (100%-specificity) that correspond to each possible cut-off value
 5 for the diagnostic test result. The cut-off value on the plot that is the closest to the upper left-hand corner (*i.e.*, the value that encloses the largest area) is the most accurate cut-off value, and a sample generating a signal that is higher than the cut-off value determined by this method may be considered positive. Alternatively, the cut-off value may be shifted to the left along the plot, to minimize the false positive rate, or to the right, to minimize the
 10 false negative rate. In general, a sample generating a signal that is higher than the cut-off value determined by this method is considered positive for a cancer.

In a related embodiment, the assay is performed in a flow-through or strip test format, wherein the binding agent is immobilized on a membrane, such as nitrocellulose. In the flow-through test, polypeptides within the sample bind to the
 15 immobilized binding agent as the sample passes through the membrane. A second, labeled binding agent then binds to the binding agent-polypeptide complex as a solution containing the second binding agent flows through the membrane. The detection of bound second binding agent may then be performed as described above. In the strip test format, one end of the membrane to which binding agent is bound is immersed in a solution containing the
 20 sample. The sample migrates along the membrane through a region containing second binding agent and to the area of immobilized binding agent. Concentration of second binding agent at the area of immobilized antibody indicates the presence of a cancer. Typically, the concentration of second binding agent at that site generates a pattern, such as a line, that can be read visually. The absence of such a pattern indicates a negative result.
 25 In general, the amount of binding agent immobilized on the membrane is selected to generate a visually discernible pattern when the biological sample contains a level of polypeptide that would be sufficient to generate a positive signal in the two-antibody sandwich assay, in the format discussed above. Preferred binding agents for use in such assays are antibodies and antigen-binding fragments thereof. Preferably, the amount of

antibody immobilized on the membrane ranges from about 25 ng to about 1 μ g, and more preferably from about 50 ng to about 500 ng. Such tests can typically be performed with a very small amount of biological sample.

Of course, numerous other assay protocols exist that are suitable for use with the tumor proteins or binding agents of the present invention. The above descriptions are intended to be exemplary only. For example, it will be apparent to those of ordinary skill in the art that the above protocols may be readily modified to use lung tumor polypeptides to detect antibodies that bind to such polypeptides in a biological sample. The detection of such lung tumor protein specific antibodies may correlate with the presence of a cancer.

A cancer may also, or alternatively, be detected based on the presence of T cells that specifically react with a lung tumor protein in a biological sample. Within certain methods, a biological sample comprising CD4⁺ and/or CD8⁺ T cells isolated from a patient is incubated with a lung tumor polypeptide, a polynucleotide encoding such a polypeptide and/or an APC that expresses at least an immunogenic portion of such a polypeptide, and the presence or absence of specific activation of the T cells is detected. Suitable biological samples include, but are not limited to, isolated T cells. For example, T cells may be isolated from a patient by routine techniques (such as by Ficoll/Hypaque density gradient centrifugation of peripheral blood lymphocytes). T cells may be incubated *in vitro* for 2-9 days (typically 4 days) at 37°C with polypeptide (*e.g.*, 5 - 25 μ g/ml). It may be desirable to incubate another aliquot of a T cell sample in the absence of lung tumor polypeptide to serve as a control. For CD4⁺ T cells, activation is preferably detected by evaluating proliferation of the T cells. For CD8⁺ T cells, activation is preferably detected by evaluating cytolytic activity. A level of proliferation that is at least two fold greater and/or a level of cytolytic activity that is at least 20% greater than in disease-free patients indicates the presence of a cancer in the patient.

As noted above, a cancer may also, or alternatively, be detected based on the level of mRNA encoding a lung tumor protein in a biological sample. For example, at least two oligonucleotide primers may be employed in a polymerase chain reaction (PCR) based

assay to amplify a portion of a lung tumor cDNA derived from a biological sample, wherein at least one of the oligonucleotide primers is specific for (*i.e.*, hybridizes to) a polynucleotide encoding the lung tumor protein. The amplified cDNA is then separated and detected using techniques well known in the art, such as gel electrophoresis. Similarly,
 5 oligonucleotide probes that specifically hybridize to a polynucleotide encoding a lung tumor protein may be used in a hybridization assay to detect the presence of polynucleotide encoding the tumor protein in a biological sample.

To permit hybridization under assay conditions, oligonucleotide primers and probes should comprise an oligonucleotide sequence that has at least about 60%, preferably
 10 at least about 75% and more preferably at least about 90%, identity to a portion of a polynucleotide encoding a lung tumor protein that is at least 10 nucleotides, and preferably at least 20 nucleotides, in length. Preferably, oligonucleotide primers and/or probes hybridize to a polynucleotide encoding a polypeptide described herein under moderately stringent conditions, as defined above. Oligonucleotide primers and/or probes which may
 15 be usefully employed in the diagnostic methods described herein preferably are at least 10-40 nucleotides in length. In a preferred embodiment, the oligonucleotide primers comprise at least 10 contiguous nucleotides, more preferably at least 15 contiguous nucleotides, of a DNA molecule having a sequence recited in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95,
 20 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 or 1676. Techniques for both PCR
 25 based assays and hybridization assays are well known in the art (*see*, for example, Mullis *et al.*, *Cold Spring Harbor Symp. Quant. Biol.*, 51:263, 1987; Erlich ed., *PCR Technology*, Stockton Press, NY, 1989).

One preferred assay employs RT-PCR, in which PCR is applied in conjunction with reverse transcription. Typically, RNA is extracted from a biological

sample, such as biopsy tissue, and is reverse transcribed to produce cDNA molecules. PCR amplification using at least one specific primer generates a cDNA molecule, which may be separated and visualized using, for example, gel electrophoresis. Amplification may be performed on biological samples taken from a test patient and from an individual who is not afflicted with a cancer. The amplification reaction may be performed on several dilutions of cDNA spanning two orders of magnitude. A two-fold or greater increase in expression in several dilutions of the test patient sample as compared to the same dilutions of the non-cancerous sample is typically considered positive.

In another embodiment, the compositions described herein may be used as markers for the progression of cancer. In this embodiment, assays as described above for the diagnosis of a cancer may be performed over time, and the change in the level of reactive polypeptide(s) or polynucleotide(s) evaluated. For example, the assays may be performed every 24-72 hours for a period of 6 months to 1 year, and thereafter performed as needed. In general, a cancer is progressing in those patients in whom the level of polypeptide or polynucleotide detected increases over time. In contrast, the cancer is not progressing when the level of reactive polypeptide or polynucleotide either remains constant or decreases with time.

Certain *in vivo* diagnostic assays may be performed directly on a tumor. One such assay involves contacting tumor cells with a binding agent. The bound binding agent may then be detected directly or indirectly via a reporter group. Such binding agents may also be used in histological applications. Alternatively, polynucleotide probes may be used within such applications.

As noted above, to improve sensitivity, multiple lung tumor protein markers may be assayed within a given sample. It will be apparent that binding agents specific for different proteins provided herein may be combined within a single assay. Further, multiple primers or probes may be used concurrently. The selection of tumor protein markers may be based on routine experiments to determine combinations that results in optimal sensitivity. In addition, or alternatively, assays for tumor proteins provided herein may be combined with assays for other known tumor antigens.

DIAGNOSTIC KITS

The present invention further provides kits for use within any of the above diagnostic methods. Such kits typically comprise two or more components necessary for performing a diagnostic assay. Components may be compounds, reagents, containers
5 and/or equipment. For example, one container within a kit may contain a monoclonal antibody or fragment thereof that specifically binds to a lung tumor protein. Such antibodies or fragments may be provided attached to a support material, as described above. One or more additional containers may enclose elements, such as reagents or buffers, to be used in the assay. Such kits may also, or alternatively, contain a detection
10 reagent as described above that contains a reporter group suitable for direct or indirect detection of antibody binding.

Alternatively, a kit may be designed to detect the level of mRNA encoding a lung tumor protein in a biological sample. Such kits generally comprise at least one oligonucleotide probe or primer, as described above, that hybridizes to a polynucleotide
15 encoding a lung tumor protein. Such an oligonucleotide may be used, for example, within a PCR or hybridization assay. Additional components that may be present within such kits include a second oligonucleotide and/or a diagnostic reagent or container to facilitate the detection of a polynucleotide encoding a lung tumor protein.

The following Examples are offered by way of illustration and not by way
20 of limitation.

EXAMPLE 1IDENTIFICATION AND CHARACTERIZATION OF LUNG
TUMOR PROTEIN cDNAS

5 This Example illustrates the identification of cDNA molecules encoding lung tumor proteins.

A. Isolation of cDNA Sequences from Lung Adenocarcinoma Libraries using Conventional cDNA Library Subtraction

10 A human lung adenocarcinoma cDNA expression library was constructed from poly A⁺ RNA from patient tissues (# 40031486) using a Superscript Plasmid System for cDNA Synthesis and Plasmid Cloning kit (BRL Life Technologies, Gaithersburg, MD) following the manufacturer's protocol. Specifically, lung carcinoma tissues were homogenized with polytron (Kinematica, Switzerland) and total RNA was extracted using Trizol reagent (BRL Life Technologies) as directed by the manufacturer. The poly A⁺ RNA was then purified using an oligo dT cellulose column as described in Sambrook et al.,
15 *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY, 1989. First-strand cDNA was synthesized using the NotI/Oligo-dT18 primer. Double-stranded cDNA was synthesized, ligated with BstXI/EcoRI adaptors (Invitrogen, San Diego, CA) and digested with NotI. Following size fractionation with cDNA size
20 fractionation columns (BRL Life Technologies), the cDNA was ligated into the BstXI/NotI site of pcDNA3.1 (Invitrogen) and transformed into ElectroMax *E. coli* DH10B cells (BRL Life Technologies) by electroporation. A total of 3 x 10⁶ independent colonies were generated.

25 Using the same procedure, a normal human cDNA expression library was prepared from a panel of normal tissue specimens, including lung, liver, pancreas, skin, kidney, brain and resting PBMC.

 cDNA library subtraction was performed using the above lung adenocarcinoma and normal tissue cDNA libraries, as described by Hara *et al.* (*Blood*,

84:189-199, 1994) with some modifications. Specifically, a lung adenocarcinoma-specific subtracted cDNA library was generated as follows. The normal tissue cDNA library (80 µg) was digested with BamHI and XhoI, followed by a filling-in reaction with DNA polymerase Klenow fragment. After phenol-chloroform extraction and ethanol precipitation, the DNA was dissolved in 133 µl of H₂O, heat-denatured and mixed with 133 µl (133 µg) of Photoprobe biotin (Vector Laboratories, Burlingame, CA). As recommended by the manufacturer, the resulting mixture was irradiated with a 270 W sunlamp on ice for 20 minutes. Additional Photoprobe biotin (67 µl) was added and the biotinylation reaction was repeated. After extraction with butanol five times, the DNA was ethanol-precipitated and dissolved in 23 µl H₂O. The resulting DNA, plus other highly redundant cDNA clones that were frequently recovered in previous lung subtractions formed the driver DNA.

To form the tracer DNA, 10 µg lung adenocarcinoma cDNA library was digested with NotI and SpeI, phenol chloroform extracted and passed through Chroma spin-400 columns (Clontech, Palo Alto, CA). Typically, 5 µg of cDNA was recovered after the sizing column. Following ethanol precipitation, the tracer DNA was dissolved in 5 µl H₂O. Tracer DNA was mixed with 15 µl driver DNA and 20 µl of 2 x hybridization buffer (1.5 M NaCl/10 mM EDTA/50 mM HEPES pH 7.5/0.2% sodium dodecyl sulfate), overlaid with mineral oil, and heat-denatured completely. The sample was immediately transferred into a 68 °C water bath and incubated for 20 hours (long hybridization [LH]). The reaction mixture was then subjected to a streptavidin treatment followed by phenol/chloroform extraction. This process was repeated three more times. Subtracted DNA was precipitated, dissolved in 12 µl H₂O, mixed with 8 µl driver DNA and 20 µl of 2 x hybridization buffer, and subjected to a hybridization at 68°C for 2 hours (short hybridization [SH]). After removal of biotinylated double-stranded DNA, subtracted cDNA was ligated into NotI/SpeI site of chloramphenicol resistant pBCSK⁺ (Stratagene, La Jolla, CA) and transformed into ElectroMax *E. coli* DH10B cells by electroporation to generate a lung adenocarcinoma specific subtracted cDNA library, referred to as LAT-S1

Similarly, LAT-S2 was generated by including 23 genes that were over-expressed in the tracer as additional drivers.

A second human lung adenocarcinoma cDNA expression library was constructed using adenocarcinoma tissue from a second patient (# 86-66) and used to
5 prepare a second lung adenocarcinoma-specific subtracted cDNA library (referred to as LAT2-S2), as described above, using the same panel of normal tissues and the additional genes over-expressed in LAT-S1.

A third human metastatic lung adenocarcinoma library was constructed from a pool of two lung pleural effusions with lung and gastric adenocarcinoma origins.
10 The subtracted cDNA library, Mets-sub2 was generated as described above using the same panel of normal tissues. However, the Mets-sub3 subtracted library was constructed by including 51 additional genes as drivers. These 51 genes were recovered in Mets-sub2, representing over-expressed housekeeping genes in the testers. As a result, Mets-sub3 is more complexed and normalized.

A total of 16 cDNA fragments isolated from LAT-S1, 585 cDNA fragments isolated from LAT-S2, 568 cDNA clones from LAT2-S2, 15 cDNA clones from Mets-sub2 and 343 cDNA clones from Mets-sub3, described above, were colony PCR amplified and their mRNA expression levels in lung tumor, normal lung, and various other normal and tumor tissues were determined using microarray technology (Incyte, Palo Alto, CA).
15 Briefly, the PCR amplification products were dotted onto slides in an array format, with each product occupying a unique location in the array. mRNA was extracted from the tissue sample to be tested, reverse transcribed, and fluorescent-labeled cDNA probes were generated. The microarrays were probed with the labeled cDNA probes, the slides scanned and fluorescence intensity was measured. This intensity correlates with the hybridization
20 intensity. Seventy-three non-redundant cDNA clones, of which 42 were found to be unique, showed over-expression in lung tumors, with expression in normal tissues tested (lung, skin, lymph node, colon, liver, pancreas, breast, heart, bone marrow, large intestine, kidney, stomach, brain, small intestine, bladder and salivary gland) being either undetectable, or at significantly lower levels compared to lung adenocarcinoma tumors.
25

These clones were further characterized by DNA sequencing with a Perkin Elmer/Applied Biosystems Division Automated Sequencer Model 373A and/or Model 377 (Foster City, CA).

The sequences were compared to known sequences in the gene bank using the EMBL GenBank databases (release 96). No significant homologies were found to the sequence provided in SEQ ID NO: 67, with no apparent homology to previously identified expressed sequence tags (ESTs). The sequences of SEQ ID NO: 60, 62, 65, 66, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97 and 98 were found to show some homology to previously identified expressed sequence tags (ESTs). The cDNA sequences of SEQ ID NO: 59, 61, 63, 64, 67, 68, 72, 73, 75, 77, 78, 81-83, 85, 87, 88, 93, 94, 96, 99 and 100 showed homology to previously identified genes. The full-length cDNA sequences for the clones of SEQ ID NO: 96 and 100 are provided in SEQ ID NO: 316 and 318, respectively. The amino acid sequences for the clones of SEQ ID NO: 59, 61, 63, 64, 68, 73, 82, 83, 94, 96 and 100 are provided in SEQ ID NO: 331, 328, 329, 332, 327, 333, 330, 326, 325, 324 and 335, respectively. A predicted amino acid sequence encoded by the sequence of SEQ ID NO: 69 (referred to as L552S) is provided in SEQ ID NO: 786.

Further studies led to the isolation of an extended cDNA sequence, and open reading frame, for L552S (SEQ ID NO: 790). The predicted amino acid sequence encoded by the cDNA sequence of SEQ ID NO: 790 is provided in SEQ ID NO: 791. The determined cDNA sequence of an isoform of L552S is provided in SEQ ID NO: 792, with the corresponding predicted amino acid sequence being provided in SEQ ID NO: 793. Subsequent studies led to the isolation of the full-length cDNA sequence of L552S (SEQ ID NO: 808). The corresponding amino acid sequence is provided in SEQ ID NO: 809. No homologies were found to the protein sequence of L552S. However, nucleotides 533-769 of the full-length cDNA sequence were found to show homology to a previously identified DNA sequence.

Full-length cloning efforts on L552S led to the isolation of three additional cDNA sequences (SEQ ID NO: 810-812) from a metastatic lung adenocarcinoma library. The sequence of SEQ ID NO: 810 was found to show some homology to previously

identified human DNA sequences. The sequence of SEQ ID NO: 811 was found to show some homology to a previously identified DNA sequence. The sequence of SEQ ID NO: 812 was found to show some homology to previously identified ESTs.

The gene of SEQ ID NO: 84 (referred to as L551S) was determined by real-time RT-PCR analysis to be over-expressed in 2/9 primary adenocarcinomas and to be expressed at lower levels in 2/2 metastatic adenocarcinomas and 1/2 squamous cell carcinomas. No expression was observed in normal tissues, with the exception of very low expression in normal stomach. Further studies on L551S led to the isolation of the 5' and 3' cDNA consensus sequences provided in SEQ ID NO: 801 and 802, respectively. The L551S 5' sequence was found to show some homology to the previously identified gene STY8 (cDNA sequence provided in SEQ ID NO: 803; corresponding amino acid sequence provided in SEQ ID NO: 805), which is a mitogen activated protein kinase phosphatase. However, no significant homologies were found to the 3' sequence of L551S. Subsequently, an extended cDNA sequence for L551S was isolated (SEQ ID NO: 804). The corresponding amino acid sequence is provided in SEQ ID NO: 806. Further studies led to the isolation of two independent full-length clones for L551S (referred to as 54298 and 54305). These two clones have five nucleotide differences compared to the STY8 DNA sequence. Two of these differences are single nucleotide polymorphisms which do not effect the encoded amino acid sequences. The other three nucleotide differences are consistent between the two L551S clones but lead to encoded amino acid sequences that are different from the STY8 protein sequence. The determined cDNA sequences for the L551S full-length clones 54305 and 54298 are provided in SEQ ID NO: 825 and 826, respectively, with the amino acid sequence for L551S being provided in SEQ ID NO: 827.

B. Isolation of cDNA Sequences from Lung Adenocarcinoma Libraries using PCR-Based cDNA Library Subtraction

cDNA clones from a PCR-based subtraction library, containing cDNA from a pool of two human lung primary adenocarcinomas subtracted against a pool of nine normal human tissue cDNAs including skin, colon, lung, esophagus, brain, kidney, spleen,

pancreas and liver, (Clontech, Palo Alto, CA) were derived and submitted to a first round of PCR amplification. This library (referred to as ALT-1) was subjected to a second round of PCR amplification, following the manufacturer's protocol. The expression levels of 760 cDNA clones in lung tumor, normal lung, and various other normal and tumor tissues, were examined using microarray technology as described above. A total of 118 clones, of which 55 were unique, were found to be over-expressed in lung tumor tissue, with expression in normal tissues tested (lung, skin, lymph node, colon, liver, pancreas, breast, heart, bone marrow, large intestine, kidney, stomach, brain, small intestine, bladder and salivary gland) being either undetectable, or at significantly lower levels. The sequences were compared to known sequences in the gene bank using the EMBL and GenBank databases (release 96). No significant homologies (including ESTs) were found to the sequence provided in SEQ ID NO: 44. The sequences of SEQ ID NO: 1, 11, 13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43, 45, 46, 51 and 57 were found to show some homology to previously identified expressed sequence tags (ESTs). The cDNA sequences of SEQ ID NO: 2-10, 12, 14, 16-19, 21, 22, 28, 31, 32, 35-38, 40, 42, 44, 47-50, 52-56 and 58 showed homology to previously identified genes. The full-length cDNA sequences for the clones of SEQ ID NO: 18, 22, 31, 35, 36 and 42 are provided in SEQ ID NO: 320, 319, 323, 321, 317, 321 and 322, respectively, with the corresponding amino acid sequences being provided in SEQ ID NO: 337, 336, 340, 338, 334, and 339, respectively.

Further studies led to the isolation of an extended cDNA sequence for the clone of SEQ ID NO: 33 (referred to as L801P). This extended cDNA sequence (provided in SEQ ID NO: 796), was found to contain three potential open reading frames (ORFs). The predicted amino acid sequences encoded by these three ORFs are provided in SEQ ID NO: 797-799, respectively. Additional full-length cloning efforts led to still further extended cDNA sequence for L801P, set forth in SEQ ID NO:1669, in addition to five potential open reading frames (ORFs 4-9; SEQ ID NOs: 1670-1675, respectively) encoded by the extended cDNA sequence. Moreover, L801P was mapped to chromosomal region 20p13 and a 137 amino acid ORF from this genomic region was identified that corresponds to ORF4 (SEQ ID NO: 1670), suggesting that this is likely an authentic ORF for L801P.

By microarray analysis, L801P was overexpressed by 2-fold or greater in the lung tumor probe groups compared to the normal tissue probe group (not shown). By real-time PCR analysis, greater than 50% of lung adenocarcinoma and greater than 30% of lung squamous cell carcinoma tumor samples tested had elevated L801P expression
 5 relative to normal lung tissue. Of those that displayed elevated L801P, the level of expression was greater than 10-fold higher than in normal lung tissue samples. Moreover, low or no expression of L801P was detected in an extensive panel of normal tissue RNAs.

We have also found that L801P expression is detected in a number of other tumor types, including breast, prostate, ovarian and colon tumors, and thus may have
 10 diagnostic and/or therapeutic utility in these cancer types as well.

In subsequent studies, a full-length cDNA sequence for the clone of SEQ ID NO: 44 (referred to as L844P) was isolated (provided in SEQ ID NO: 800). Comparison of this sequence with those in the public databases revealed that the 470 bases at the 5' end of the sequence show homology to the known gene dihydrodiol dehydrogenase, thus
 15 indicating that L844P is a novel transcript of the dihydrodiol dehydrogenase family having 2007 base pairs of previously unidentified 3' untranslated region.

The predicted amino acid sequence encoded by the sequence of SEQ ID NO: 46 (referred to as L840P) is provided in SEQ ID NO: 787. An extended cDNA sequence for L840P, which was determined to include an open reading frame, is provided
 20 in SEQ ID NO: 794. The predicted amino acid sequence encoded by the cDNA sequence of SEQ ID NO: 794 is provided in SEQ ID NO: 795. The full-length cDNA sequence for the clone of SEQ ID NO: 54 (referred to as L548S) is provided in SEQ ID NO: 788, with the corresponding amino acid sequence being provided in SEQ ID NO: 789.

Northern blot analyses of the genes of SEQ ID NO: 25 and 46 (referred to as
 25 L839P and L840P, respectively) were remarkably similar. Both genes were expressed in 1/2 lung adenocarcinomas as two bands of 3.6 kb and 1.6 kb. No expression of L839P was observed in normal lung or trachea. No expression of L840P was observed in normal bone marrow, resting or activated PBMC, esophagus, or normal lung. Given the similar expression patterns, L839P and L840P may be derived from the same gene.

Further studies on L773P (SEQ ID NO: 58) resulted in the isolation of the extended consensus cDNA sequence provided in SEQ ID NO: 807.

Additional lung adenocarcinoma cDNA clones were isolated as follows. A cDNA library was prepared from a pool of two lung adenocarcinomas and subtracted
 5 against cDNA from a panel of normal tissues including lung, brain, liver, kidney, pancreas, skin, heart and spleen. The subtraction was performed using a PCR-based protocol (Clontech), which was modified to generate larger fragments. Within this protocol, tester and driver double stranded cDNA were separately digested with five restriction enzymes that recognize six-nucleotide restriction sites (MluI, MscI, PvuII, SalI and StuI). This
 10 digestion resulted in an average cDNA size of 600 bp, rather than the average size of 300 bp that results from digestion with RsaI according to the Clontech protocol. The ends of the restriction digested tester cDNA were filled in to generate blunt ends for adapter ligation. This modification did not affect the subtraction efficiency. Two tester populations were then created with different adapters, and the driver library remained
 15 without adapters. The tester and driver libraries were then hybridized using excess driver cDNA. In the first hybridization step, driver was separately hybridized with each of the two tester cDNA populations. This resulted in populations of (a) unhybridized tester cDNAs, (b) tester cDNAs hybridized to other tester cDNAs, (c) tester cDNAs hybridized to driver cDNAs and (d) unhybridized driver cDNAs. The two separate hybridization
 20 reactions were then combined, and rehybridized in the presence of additional denatured driver cDNA. Following this second hybridization, in addition to populations (a) through (d), a fifth population (e) was generated in which tester cDNA with one adapter hybridized to tester cDNA with the second adapter. Accordingly, the second hybridization step resulted in enrichment of differentially expressed sequences which could be used as
 25 templates for PCR amplification with adaptor-specific primers.

The ends were then filled in, and PCR amplification was performed using adaptor-specific primers. Only population (e), which contained tester cDNA that did not hybridize to driver cDNA, was amplified exponentially. A second PCR amplification step

was then performed, to reduce background and further enrich differentially expressed sequences.

Fifty-seven cDNA clones were isolated from the subtracted library (referred to as LAP1) and sequenced. The determined cDNA sequences for 16 of these clones are provided in SEQ ID NO: 101-116. The sequences of SEQ ID NO: 101 and 114 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 102-109 and 112 showed some similarity to previously identified sequences, while the sequences of SEQ ID NO: 113, 115 and 116 showed some similarity to previously isolated ESTs.

An additional 502 clones analyzed from the LAP1 library were sequenced and the determined cDNA sequences are shown in SEQ ID NO:828-1239 and 1564-1653.

C. Isolation of cDNA Sequences from Small Cell Lung Carcinoma Libraries using PCR-Based cDNA Library Subtraction

A subtracted cDNA library for small cell lung carcinoma (referred to as SCL1) was prepared using essentially the modified PCR-based subtraction process described above. cDNA from small cell lung carcinoma was subtracted against cDNA from a panel of normal tissues, including normal lung, brain, kidney, liver, pancreas, skin, heart, lymph node and spleen. Both tester and driver poly A⁺ RNA were initially amplified using SMART PCR cDNA synthesis kit (Clontech, Palo Alto, CA). The tester and driver double stranded cDNA were separately digested with five restriction enzymes (DraI, MscI, PvuII, SmaI, and StuI). These restriction enzymes generated blunt end cuts and the digestion resulted in an average insert size of 600 bp. Digestion with this set of restriction enzymes eliminates the step required to generate blunt ends by filling in of the cDNA ends. These modifications did not affect subtraction efficiency.

Eighty-five clones were isolated and sequenced. The determined cDNA sequences for 31 of these clones are provided in SEQ ID NO: 117-147. The sequences of SEQ ID NO: 122, 124, 126, 127, 130, 131, 133, 136, 139 and 147 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 120, 129,

135, 137, 140, 142, 144 and 145 showed some similarity to previously identified gene sequences, while the sequences of SEQ ID NO: 114, 118, 119, 121, 123, 125, 128, 132, 134, 138, 141, 143 and 147 showed some similarity to previously isolated ESTs.

In further studies, three additional cDNA libraries were generated from poly A+ RNA from a single small cell lung carcinoma sample subtracted against a pool of poly A+ RNA from nine normal tissues (lung, brain, kidney, liver, pancreas, skin, heart pituitary gland and spleen). For the first library (referred to as SCL2), the subtraction was carried out essentially as described above for the LAP1 library, with the exception that the tester and driver were digested with PvuII, StuI, MscI and DraI. The ratio of tester and driver cDNA used was as recommended by Clontech. For the second library (referred to as SCL3), subtraction was performed essentially as for SCL2 except that cDNA for highly redundant clones identified from the SCL2 library was included in the driver cDNA. Construction of the SCL4 library was performed essentially as described for the SCL3 library except that a higher ratio of driver to tester was employed.

Each library was characterized by DNA sequencing and database analyses. The determined cDNA sequence for 35 clones isolated from the SCL2 library are provided in SEQ ID NO: 245-279, with the determined cDNA sequences for 21 clones isolated from the SCL3 library and for 15 clones isolated from the SCL4 library being provided in SEQ ID NO: 280-300 and 301-315, respectively. The sequences of SEQ ID NO: 246, 254, 261, 262, 304, 309 and 311 showed no significant homologies to previously identified sequences. The sequence of SEQ ID NO: 245, 248, 255, 266, 270, 275, 280, 282, 283, 288-290, 292, 295, 301 and 303 showed some homology to previously isolated ESTs, while the sequences of SEQ ID NO: 247, 249-253, 256-260, 263-265, 267-269, 271-274, 276-279, 281, 284-287, 291, 293, 294, 296-300, 302, 305-308, 310 and 312-315 showed some homology to previously identified gene sequences.

3264 cDNA clones from three PCR-based subtracted cDNA libraries were analyzed by cDNA microarray technology as part of Lung Chip 5. Of the 3264 cDNA clones 960 clones came from SQL1 library, 768 clones came from SCL1 library, and 1536 clones came from SCL3 and SCL4 libraries. 35 pairs of fluorescent labeled cDNA probes

were used for the microarray analysis. Each probe pair included a lung tumor probe paired with a normal tissue probe. The expression data was analyzed. 498 cDNA clones were found to be overexpressed by 2-fold or greater in the small cell and/or non-small cell lung tumor probe groups compared to the normal tissue probe group. Also, the mean expression values for these clones in normal tissues were below 0.1 (range of expression is from 0.001 to 10). The cDNA sequences disclosed in SEQ ID NO:1240-1563 represent 324 non-redundant clones.

The following sequences were novel based on database analysis including GenBank and GeneSeq: SEQ ID NO:1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, and 1563.

Full-length sequence for contig 139 (SEQ ID NO: 1467), also known as L985P, was identified by searching public databases using SEQ ID NO: 1467 as a query. By this approach, L985 was identified as cell surface immunomodulator-2 (CSIMM-2), the cDNA sequence of which is set forth in SEQ ID NO: 1676, encoding a protein having the sequence set forth in SEQ ID NO: 1677.

By microarray analysis, L985P was overexpressed by 2-fold or greater in the lung tumor probe groups compared to the normal tissue probe group. Moreover, the mean expression values for L985P in normal tissues was below 0.2 (range of expression was from 0.01 to 10). By real-time PCR analysis, greater than 40% of small cell lung carcinoma lung tumor samples tested had elevated L985P expression relative to normal lung tissue. Of those that displayed elevated L985P, the level of expression was greater than 3-fold higher than in normal lung tissue samples. Low or no expression of L985P was detected in an extensive panel of normal tissue RNAs. These findings for L985P support its use both as a diagnostic marker for detecting the presence of lung cancer in a patient and/or as an immunotherapeutic target for the treatment of lung cancer.

D. Isolation of cDNA Sequences from a Neuroendocrine Library using PCR-Based cDNA Library Subtraction

Using the modified PCR-based subtraction process, essentially as described above for the LAP1 subtracted library, a subtracted cDNA library (referred to as MLN1) was derived from a lung neuroendocrine carcinoma that had metastasized to the subcarinal lymph node, by subtraction with a panel of nine normal tissues, including normal lung, brain, kidney, liver, pancreas, skin, heart, lymph node and spleen.

Ninety-one individual clones were isolated and sequenced. The determined cDNA sequences for 58 of these clones are provided in SEQ ID NO: 147-222. The sequences of SEQ ID NO: 150, 151, 154, 157, 158, 159, 160, 163, 174, 175, 178, 186-190, 192, 193, 195-200, 208-210, 212-215 and 220 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 152, 155, 156, 161, 165, 166, 176, 179, 182, 184, 185, 191, 194, 221 and 222 showed some similarity to previously identified gene sequences, while the sequences of SEQ ID NO: 148, 149, 153, 164, 167-173, 177, 180, 181, 183, 201-207, 211 and 216-219 showed some similarity to previously isolated ESTs.

The determined cDNA sequences of an additional 442 clones isolated from the MLN1 library are provided in SEQ ID NO: 341-782. The determined cDNA sequences of an additional 11 clones isolated from the MLN1 library are provided in SEQ ID NO:1654-1664.

E. Isolation of cDNA Sequences from a Squamous Cell Lung Carcinoma Library using PCR-Based cDNA Library Subtraction

A subtracted cDNA library for squamous cell lung carcinoma (referred to as SQL1) was prepared, essentially using the modified PCR-based subtraction process described above, except the tester and driver double stranded cDNA were separately digested with four restriction enzymes (DraI, MscI, PvuII and StuI) cDNA from a pool of two squamous cell lung carcinomas was subtracted against cDNA from a pool of 10 normal

tissues, including normal lung, brain, kidney, liver, pancreas, skin, heart, spleen, esophagus and trachea.

Seventy-four clones were isolated and sequenced. The determined cDNA sequences for 22 of these clones are provided in SEQ ID NO: 223-244. The sequence of
 5 SEQ ID NO: 241 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 223, 225, 232, 233, 235, 238, 239, 242 and 243 showed some similarity to previously identified gene sequences, while the sequences of SEQ ID NO: 224, 226-231, 234, 236, 237, 240, 241 and 244 showed some similarity to previously isolated ESTs.

10 The sequences of an additional 12 clones isolated during characterization of cDNA libraries prepared from lung tumor tissue are provided in SEQ ID NO: 813-824. Comparison of these sequences with those in the GenBank database and the GeneSeq DNA database revealed no significant homologies to previously identified sequences.

15

EXAMPLE 2

SYNTHESIS OF POLYPEPTIDES

Polypeptides may be synthesized on a Perkin Elmer/Applied Biosystems Division 430A peptide synthesizer using FMOC chemistry with HPTU (O-Benzotriazole-
 20 N,N,N',N'-tetramethyluronium hexafluorophosphate) activation. A Gly-Cys-Gly sequence may be attached to the amino terminus of the peptide to provide a method of conjugation, binding to an immobilized surface, or labeling of the peptide. Cleavage of the peptides from the solid support may be carried out using the following cleavage mixture: trifluoroacetic acid:ethanedithiol:thioanisole:water :phenol (40:1:2:2:3). After cleaving for
 25 2 hours, the peptides may be precipitated in cold methyl-t-butyl-ether. The peptide pellets may then be dissolved in water containing 0.1% trifluoroacetic acid (TFA) and lyophilized prior to purification by C18 reverse phase HPLC. A gradient of 0%-60% acetonitrile (containing 0.1% TFA) in water (containing 0.1% TFA) may be used to elute the peptides.

Following lyophilization of the pure fractions, the peptides may be characterized using electrospray or other types of mass spectrometry and by amino acid analysis.

EXAMPLE 3

5 PREPARATION OF ANTIBODIES AGAINST LUNG CANCER ANTIGENS

Polyclonal antibodies against the lung cancer antigen L773P (SEQ ID NO: 783) were prepared as follows.

Rabbits were immunized with recombinant protein expressed in and purified
10 from *E. coli* as described above. For the initial immunization, 400 µg of antigen combined with muramyl dipeptide (MDP) was injected subcutaneously (S.C.). Animals were boosted S.C. 4 weeks later with 200 µg of antigen mixed with incomplete Freund's Adjuvant (IFA). Subsequent boosts of 100 µg of antigen mixed with IFA were injected S.C. as necessary to induce high antibody titer responses. Serum bleeds from immunized rabbits were tested for
15 L773P-specific reactivity using ELISA assays with purified protein and showed strong reactivity to L773P. Polyclonal antibodies against L773P were affinity purified from high titer polyclonal sera using purified protein attached to a solid support.

EXAMPLE 4

20 PROTEIN EXPRESSION OF LUNG TUMOR-SPECIFIC ANTIGENS

Full-length L773P (amino acids 2-364 of SEQ ID NO: 783), with a 6X His Tag, were subcloned into the pPDM expression vector and transformed into either BL21 CodonPlus or BL21 pLysS host cells using standard techniques. High levels of expression
25 were observed in both cases. Similarly, the N-terminal portion of L773P (amino acids 2-71 of SEQ ID NO: 783; referred to as L773PA), with a 6X His tag were subcloned into the vector pPDM and transformed into BL21 CodonPlus host cells. Low levels of expression were observed by N-terminal sequencing. The sequence of the expressed constructs for L773P and L773PA are provided in SEQ ID NO: 784 and 785, respectively.

EXAMPLE 5

EXPRESSION IN E. COLI OF L548S HIS TAG FUSION PROTEIN

5 The L548S coding region was PCR amplified with the following primers:

Forward primer starting at amino acid 2:

PDM-433: 5' gctaaaggtgaccccaagaaccaaag 3' Tm 60°C (SEQ ID NO:1665)

Reverse primer creating a XhoI site after the stop codon:

10 PDM-438: 5' ctattaactcgagggagacagataaacagtttcttta 3' Tm 61°C (SEQ ID NO:1666)

 The PCR product was then digested with XhoI restriction enzyme, gel purified and then cloned into pPDM His, a modified pET28 vector with a His tag in frame, which had been digested with Eco72I and XhoI restriction enzymes. The correct construct
15 was confirmed by DNA sequence analysis and then transformed into BL21 (DE3) pLys S and BL21 (DE3) CodonPlus RIL expression hosts.

 The protein sequence of expressed recombinant L548S is shown in SEQ ID NO:1667, and the DNA sequence of expressed recombinant L7548S is shown in
20 SEQ ID NO:1668.

 From the foregoing it will be appreciated that, although specific embodiments of the invention have been described herein for purposes of illustration, various modifications may be made without deviating from the spirit and scope of the
25 invention. Accordingly, the invention is not limited except as by the appended claims.

CLAIMS

What is claimed:

1. An isolated polypeptide, comprising at least an immunogenic portion of a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669;

(b) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 under moderately stringent conditions; and

(c) complements of sequences of (a) or (b).

2. An isolated polypeptide according to claim 1, wherein the polypeptide comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, and 1669 or a complement of any of the foregoing polynucleotide sequences.

3. An isolated polypeptide comprising a sequence recited in any one of SEQ ID NOs: 786, 787, 791, 793, 795, 797-799, 806, 809, 827 and 1670-1675.

4. An isolated polynucleotide encoding at least 15 amino acid residues of a lung tumor protein, or a variant thereof that differs in one or more substitutions, deletions, additions and/or insertions such that the ability of the variant to react with antigen-specific antisera is not substantially diminished, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide comprising a sequence recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785,

790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 or a complement of any of the foregoing sequences.

5. An isolated polynucleotide encoding a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide comprising a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 or a complement of any of the foregoing sequences.

6. An isolated polynucleotide, comprising a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280,

1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669.

7. An isolated polynucleotide, comprising a sequence that hybridizes to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 under moderately stringent conditions.

8. An isolated polynucleotide complementary to a polynucleotide according to any one of claims 4-7.

9. An expression vector, comprising a polynucleotide according to any one of claims 4-8.

10. A host cell transformed or transfected with an expression vector according to claim 9.

11. An isolated antibody, or antigen-binding fragment thereof, that specifically binds to a lung tumor protein that comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 or a complement of any of the foregoing polynucleotide sequences.

12. A fusion protein, comprising at least one polypeptide according to claim 1.

13. A fusion protein according to claim 12, wherein the fusion protein comprises an expression enhancer that increases expression of the fusion protein in a host cell transfected with a polynucleotide encoding the fusion protein.

14. A fusion protein according to claim 12, wherein the fusion protein comprises a T helper epitope that is not present within the polypeptide of claim 1.

15. A fusion protein according to claim 12, wherein the fusion protein comprises an affinity tag.

16. An isolated polynucleotide encoding a fusion protein according to claim 12.

17. A pharmaceutical composition, comprising a physiologically acceptable carrier and at least one component selected from the group consisting of:

- (a) a polypeptide according to claim 1;
- (b) a polynucleotide according to claim 4;
- (c) an antibody according to claim 11;
- (d) a fusion protein according to claim 12; and
- (e) a polynucleotide according to claim 16.

18. An immunogenic composition comprising an immunostimulant and at least one component selected from the group consisting of:

- (a) a polypeptide according to claim 1;
- (b) a polynucleotide according to claim 4;
- (c) an antibody according to claim 11;
- (d) a fusion protein according to claim 12; and
- (e) a polynucleotide according to claim 16.

19. An immunogenic composition according to claim 18, wherein the immunostimulant is an adjuvant.

20. An immunogenic composition according to any claim 18, wherein the immunostimulant induces a predominantly Type I response.

21. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a pharmaceutical composition according to claim 17.

22. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of an immunogenic composition according to claim 18.

23. A pharmaceutical composition comprising an antigen-presenting cell that expresses a polypeptide according to claim 1, in combination with a pharmaceutically acceptable carrier or excipient.

24. A pharmaceutical composition according to claim 23, wherein the antigen presenting cell is a dendritic cell or a macrophage.

25. An immunogenic composition comprising an antigen-presenting cell that expresses a polypeptide comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 and 828-1664, 1669 and 1676;

(b) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 under moderately stringent conditions; and

(c) complements of sequences of (i) or (ii);

in combination with an immunostimulant.

26. An immunogenic composition according to claim 25, wherein the immunostimulant is an adjuvant.

27. An immunogenic composition according to claim 25, wherein the immunostimulant induces a predominantly Type I response.

28. An immunogenic composition according to claim 25, wherein the antigen-presenting cell is a dendritic cell.

29. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of an antigen-presenting cell that expresses a polypeptide comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676;

(b) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676 under moderately stringent conditions; and

(c) complements of sequences of (i) or (ii) encoded by a polynucleotide recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676;

and thereby inhibiting the development of a cancer in the patient.

30. A method according to claim 29, wherein the antigen-presenting cell is a dendritic cell.

31. A method according to any one of claims 21, 22 and 29, wherein the cancer is lung cancer.

32. A method for removing tumor cells from a biological sample, comprising contacting a biological sample with T cells that specifically react with a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(i) polynucleotides recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676 ; and

(ii) complements of the foregoing polynucleotides;

wherein the step of contacting is performed under conditions and for a time sufficient to permit the removal of cells expressing the antigen from the sample.

33. A method according to claim 32, wherein the biological sample is blood or a fraction thereof.

34. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient a biological sample treated according to the method of claim 32.

35. A method for stimulating and/or expanding T cells specific for a lung tumor protein, comprising contacting T cells with at least one component selected from the group consisting of:

(a) polypeptides comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(i) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676;

(ii) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785,

790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676 under moderately stringent conditions; and

- (iii) complements of sequences of (i) or (ii);
- (b) polynucleotides encoding a polypeptide of (a); and
- (c) antigen presenting cells that express a polypeptide of (a);

under conditions and for a time sufficient to permit the stimulation and/or expansion of T cells.

36. An isolated T cell population, comprising T cells prepared according to the method of claim 35.

37. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a T cell population according to claim 36.

38. A method for inhibiting the development of a cancer in a patient, comprising the steps of:

(a) incubating CD4⁺ and/or CD8⁺ T cells isolated from a patient with at least one component selected from the group consisting of:

(i) polypeptides comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(1) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676;

(2) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 under moderately stringent conditions; and

(3) complements of sequences of (1) or (2);

(ii) polynucleotides encoding a polypeptide of (i); and

(iii) antigen presenting cells that expresses a polypeptide of (i);

such that T cells proliferate; and

(b) administering to the patient an effective amount of the proliferated T cells, and thereby inhibiting the development of a cancer in the patient.

39. A method for inhibiting the development of a cancer in a patient, comprising the steps of:

(a) incubating CD4⁺ and/or CD8⁺ T cells isolated from a patient with at least one component selected from the group consisting of:

(i) polypeptides comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(1) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676;

(2) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 under moderately stringent conditions; and

(3) complements of sequences of (1) or (2);

(ii) polynucleotides encoding a polypeptide of (i); and

(iii) antigen presenting cells that express a polypeptide of (i);

such that T cells proliferate;

(b) cloning at least one proliferated cell to provide cloned T cells; and

(c) administering to the patient an effective amount of the cloned T cells, and thereby inhibiting the development of a cancer in the patient.

40. A method for determining the presence or absence of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with a binding agent that binds to a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of polypeptide that binds to the binding agent; and

(c) comparing the amount of polypeptide to a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient.

41. A method according to claim 40, wherein the binding agent is an antibody.

42. A method according to claim 43, wherein the antibody is a monoclonal antibody.

43. A method according to claim 40, wherein the cancer is lung cancer.

44. A method for monitoring the progression of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient at a first point in time with a binding agent that binds to a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of polypeptide that binds to the binding agent;

(c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and

(d) comparing the amount of polypeptide detected in step (c) to the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

45. A method according to claim 44, wherein the binding agent is an antibody.

46. A method according to claim 45, wherein the antibody is a monoclonal antibody.

47. A method according to claim 44, wherein the cancer is a lung cancer.

48. A method for determining the presence or absence of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide; and

(c) comparing the amount of polynucleotide that hybridizes to the oligonucleotide to a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient.

49. A method according to claim 48, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a polymerase chain reaction.

50. A method according to claim 48, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a hybridization assay.

51. A method for monitoring the progression of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide;

(c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and

(d) comparing the amount of polynucleotide detected in step (c) to the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

52. A method according to claim 51, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a polymerase chain reaction.

53. A method according to claim 51, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a hybridization assay.

54. A diagnostic kit, comprising:

- (a) one or more antibodies according to claim 11; and
- (b) a detection reagent comprising a reporter group.

55. A kit according to claim 54, wherein the antibodies are immobilized on a solid support.

56. A kit according to claim 54, wherein the detection reagent comprises an anti-immunoglobulin, protein G, protein A or lectin.

57. A kit according to claim 54, wherein the reporter group is selected from the group consisting of radioisotopes, fluorescent groups, luminescent groups, enzymes, biotin and dye particles.

58. An oligonucleotide comprising 10 to 40 contiguous nucleotides that hybridize under moderately stringent conditions to a polynucleotide that encodes a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475,

1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, 1669 and 1676 or a complement of any of the foregoing polynucleotides.

59. A oligonucleotide according to claim 58, wherein the oligonucleotide comprises 10-40 contiguous nucleotides recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, 1669 and 1676.

60. A diagnostic kit, comprising:

- (a) an oligonucleotide according to claim 59; and
- (b) a diagnostic reagent for use in a polymerase chain reaction or hybridization assay.

COMPOSITIONS AND METHODS FOR THE
THERAPY AND DIAGNOSIS OF LUNG CANCER

ABSTRACT OF THE DISCLOSURE

Compositions and methods for the therapy and diagnosis of cancer, such as lung cancer, are disclosed. Compositions may comprise one or more lung tumor proteins, immunogenic portions thereof, or polynucleotides that encode such portions. Alternatively, a therapeutic composition may comprise an antigen presenting cell that expresses a lung tumor protein, or a T cell that is specific for cells expressing such a protein. Such compositions may be used, for example, for the prevention and treatment of diseases such as lung cancer. Diagnostic methods based on detecting a lung tumor protein, or mRNA encoding such a protein, in a sample are also provided.

WPN\210121-Corixa\478c10\478c10-app.doc

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants : Tongtong Wang et al.
Filed : August 29, 2000
For : COMPOSITIONS AND METHODS FOR ^{THE} THERAPY AND
DIAGNOSIS OF LUNG CANCER

Docket No. : 210121.478C10

Date : August 29, 2000

Box Patent Application
Assistant Commissioner for Patents
Washington, D.C. 20231

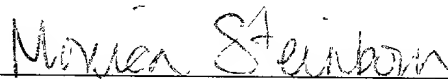
DECLARATION

Sir:

I, Monica Steinborn, in accordance with 37 C.F.R. § 1.821(f) do hereby declare that, to the best of my knowledge, the content of the paper entitled "Sequence Listing" and the computer readable copy contained within the floppy disk are the same.

I declare further that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true and further that these statements are made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issuing thereon.

Dated this 29th day of August, 2000.



Monica Steinborn
Biotechnology Paralegal

701 Fifth Avenue, Suite 6300
Seattle, WA 98104-7092
(206) 622-4900
FAX (206) 682-6031

006280 "E3T560"

SEQUENCE LISTING

<110> Wang, Tongtong
 Bangur, Chaitanya S.
 Lodes, Michael A.
 Fanger, Gary
 Vedvick, Tom
 Carter, Darrick
 Retter, Marc
 Mannion, Jane
 Fan, Liqun

<120> COMPOSITIONS AND METHODS FOR ^{THE} THERAPY AND
 DIAGNOSIS OF LUNG CANCER

<130> 210121.478C10

<140> US

<141> 2000-08-29

<160> 1679

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 527

<212> DNA

<213> Homo sapien

<400> 1

| | | | | | | |
|-------------|------------|-------------|------------|------------|-------------|-----|
| ccaccagtcc | acaaatgtga | ctggtaaggg | atctagtaac | agaggatgga | gttgggcaga | 60 |
| atattatcct | ggatgatatg | caccagcac | tagaatacac | ctttcattag | aatgaagaga | 120 |
| acagacaaag | ccctcagaaa | agatacaaag | gcagagacat | tgattagaac | attatctcat | 180 |
| aacagagggtg | gggccattac | ccaccattat | tgtaaaataa | ctgtaactaa | ccaaaacaca | 240 |
| tacaggcttc | tttaatggag | ttaataaaaac | tatggcacat | tgggaatcag | gggcagagggt | 300 |
| actgttccca | gacggaaaac | tgggataaag | ggagccatgc | tgacagggcc | ttattccagt | 360 |
| ctaggttggt | agaaaggagc | cctagcccag | aaatgacagc | aaatagccat | aatcattatg | 420 |
| tggggctgaa | ccagaggaag | ccaggctgag | ccaagaagct | ggaagtatct | tgaacggctc | 480 |
| tccaaatcca | aagattatcc | atactcttta | tccctccagc | gatgtgt | | 527 |

<210> 2

<211> 490

<212> DNA

<213> Homo sapien

<400> 2

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaagagttc | tccactgtga | agactgaaag | gacctggtga | catttcggca | tcagtcctgt | 60 |
| taccacttgg | aggtaacaga | agcaggctcg | tgctctcctt | taattctacc | acactacatg | 120 |
| actcgcaatt | ggttctgaaa | ttagaacgtt | caccatcgta | cttaaaatct | taggggcatg | 180 |
| aagagtcagc | tagaacaagg | aaaaagaaag | tcgcaggtag | taggtaagta | ggtgggcaca | 240 |
| tgaaaagcca | agctgctctg | tccaacacca | gtgtacatgt | gctttaacta | aatgaactcc | 300 |
| agaggccaac | agcagcagac | ctgctcaatt | caccttccaa | atcagaacaa | gacccaaaag | 360 |
| ctcaggcttg | agttgtcaac | tatgcatagg | ttccgccagt | gctgaggggt | gtgaggctct | 420 |
| agttgtgaag | aagctacaag | aaatcatgat | gcatgtgatc | tgggccgcac | tggcatttgc | 480 |
| agctattcag | | | | | | 490 |

<210> 3
 <211> 464
 <212> DNA
 <213> Homo sapien

<400> 3
 ggagctgtgg gctcagtcgt ggggcagatt gcaaagctca agggctgcaa agttgttggg 60
 gcagtagggg ctgatgaaaa gggtgcctac cttcaaaagc ttggatttga tgtcgtcttt 120
 aactacaaga cggtagagtc tttggaagaa accttgaaga aagcgtctcc tgatggttat 180
 gattgttatt ttgataatgt aggtggagag ttttcaaaca ctgttatcgg ccagatgaag 240
 aaatttggaa ggattgccat atgtggagcc atctctacat ataacagaac cggcccactt 300
 cccccaggcc cccccccaga gattgttatc tatcaggagc ttcgcatgga agcttttgtc 360
 gtctaccgct ggcaaggaga tgcccgccaa aaagctctga aggacttgct gaaatgggtc 420
 ttagagttta aatttcagct tccctacttt gtaattgact gact 464

<210> 4
 <211> 510
 <212> DNA
 <213> Homo sapien

<400> 4
 ccttatcaca ctgtaagtgg tccaagccca tagggatgct ctttttggtt cctggaattt 60
 ccagttggat gtgacagaga tctttcagta taggtctaag tcaagagtag cctctggggt 120
 gaggtgggct gggagattaa catcttacct ggggtccttc agataaacct gttggttttt 180
 cctgtctcat acaggcccat cttaagtttt gatgttgaat taaaactact tctaccccct 240
 tagttataaa aaaggccaca aggagcattt atgtggatat ctggaagtga gatagttatt 300
 ccattcccag gaaaagaaaa ataaagctaa gttacaaaac taaatctata tgcaataaag 360
 ttattatata ctgctttgtt taagcagagt cctctggaat ttatgtacag tacattagtt 420
 ttcagctatt tatattccac aagttagacc ttaagattct ctgggtttta gacaattgtt 480
 aaagatactt ctaaagctct gaggcagttca 510

<210> 5
 <211> 452
 <212> DNA
 <213> Homo sapien

<400> 5
 acagcgcctc acgcacctga gccccgagga gaaggcgtcg aggaggaaac tgaaaaacag 60
 agtagcagct cagactgcca gagatcgaaa gaaggctcga atgagtgagc tggaacagca 120
 agtggtagat ttagaagaag agaaccaaaa acttttgcta gaaaatcagc ttttacgaga 180
 gaaaactcat ggccttgtag ttgagaacca ggagttaaga cagcgttgg ggatggatgc 240
 cctggttgct gaagaggagg cggaagccaa ggtaaatcat ctcttttatt tgggtgectca 300
 tgtgagtact ggttccaagt gacatgaccc agcgattatg tttacagtct ggactttctga 360
 tcaagagcgt tcttgaaatt ttccttcagt ttttaagacat tttcatgcag gcagagtgtt 420
 cttcccctaa aggcacttga cactcatttt tt 452

<210> 6
 <211> 336
 <212> DNA
 <213> Homo sapien

<400> 6
 tatagagtgc tgacatctga cattgagaaa ttcatgccta ttgtttatac tcccactgtg 60

```

ggctctggctt gccacaata tagtttggtg tttcggaagc caagaggtct ctttattact 120
atccacgatc gagggcatat tgcttcagtt ctcaatgcac ggccagaaga tgtcatcaag 180
gccattgtgg tgactgatgg agagcgtatt cttggcttgg gagaccttgg ctgtaatgga 240
atgggcatcc ctgtgggtaa attggctcta tatacagctt gcggagggat gaatcctcaa 300
gaatgtctgc ctgtcattct ggatgtggga accgaa 336

```

```

<210> 7
<211> 376
<212> DNA
<213> Homo sapien

```

```

<400> 7
ctgtgggaaa cctcattgtt ctgtacaaag tactagctaa accagaaagg tgattccagg 60
aggagtttagc caaacaacaa caaaaacaaa aaatgtgctg ttcaagtttt cagctttaag 120
atatctttgg ataattgttat ttctatTTTT tatttttttt cattagaagt taccaaatta 180
agatggtaag acctctgaga ccaaaatTTT gtcccatctc taccctctca caactgctta 240
cagaatggat catgtcccc ttatgttgag gtgaccactt aattgctttc ctgcctcctt 300
gaaagaaaga aagaaagaag actgtgtttt tgccactgat ttagccatgt gaaactcatc 360
tcattaccct tttctg 376

```

```

<210> 8
<211> 406
<212> DNA
<213> Homo sapien

```

```

<400> 8
ggtagggagc aattctatta tttggcattg catggctggg ttgaattaaa acagggagtg 60
agaacaggtg agtctagaag tccaactctg aaaaggacca ctgtacattt gaacacacgg 120
ctgtgttaaa gatgctgcta atgtcagtcac ctgggtgcac taaaggatct cttatTTTtat 180
gtaaaacggt gggattgaca agatagatct gatactctgt taagttaccc tctgaagcta 240
cttcttgtga aataactaat acagcatcat cctgccaaagc gaaagaggca ggcataagca 300
aggacaaatt aaaagggggg aagagcctta tcatgatgag gagtcttgtt ttgacatctt 360
gggaaaagct gtccatagtg tgaagtcgtc aatttctcac catggt 406

```

```

<210> 9
<211> 330
<212> DNA
<213> Homo sapien

```

```

<400> 9
actactacca agagctgcag agagacattt ctgaaatgtt tttgcagatt tataaacaag 60
ggggTTTTct gggcctctcc aatattaagt tcaggccagg atctgtgggtg gtacaattga 120
ctctggcctt ccgagaagg accatcaatg tccacgacgt ggagacacag ttcaatcagt 180
ataaaacgga agcagcctct cgatataacc tgacgatctc agacgtcagc gtgagtgatg 240
tgccatttcc tttctctgcc cagtctgggg ctgggggtgcc aggctggggc atcgcgctgc 300
tgggtgctgg ctgtgttctg gttgcgctgg 330

```

```

<210> 10
<211> 449
<212> DNA
<213> Homo sapien

```

```

<400> 10
ctgacggctt tgctgtccca gagccgcta aacgcaagaa aagtcgatgg gacagttaga 60

```

```

ggggatgtgc taaagcgtga aatcagttgt ccttaatttt tagaaagatt ttggtaacta 120
ggtgtctcag ggctgggttg gggccaaaag tgtaaggacc ccctgccctt agtggagagc 180
tgagagcttg agacattacc ccttcacagc aaggaatttt cggatgtttt cttgggaagc 240
tgttttggtc cttggaagca gtgagagctg ggaagcttct tttggctcta ggtgagttgt 300
catgcgggta agttgaggtt atcttgggat aaagggtctt ctagggcaca aaactcactc 360
taggtttata ttgtatgtag cttatatttt ttactaaggt gtcaccttat aagcatctat 420
aaattgagtt ctttttctta gttgtatgg 449

```

```

<210> 11
<211> 472
<212> DNA
<213> Homo sapien

```

```

<400> 11
cctcgatgca tgctgctcta cctctcatca gccacagtc tgacacgagg tcctctttgg 60
tctgtggtga ggtatggatg tctgcagtct acacaacagc cctgcagaac gggcctggac 120
aacccttggg ggataagaca gccacacatg gctcaggctg ttaggtgtcc actgtcacag 180
tccaaagaga aaggtagcggc ctccaagggg gcagcttaag ccaacatgta agacttgggc 240
acgatgaaaag gacggggggtc cagctacgaa tgtttttggc cttgatgtca agttgccagc 300
tactggaagg caggagcagt ttctttcttt tcccactctg tgctgggtac ttgggagagg 360
cgaaataaat accagactgt ccaactcctca gcctaaggtc cttctcaagt cctgcacact 420
cagcacttgc tctttaacgt ggcataatgt ccccatctt cccctggtta tg 472

```

```

<210> 12
<211> 371
<212> DNA
<213> Homo sapien

```

```

<400> 12
tttttttttt tttttttttt ttttggarat ttgkacatt ttattcagwa tttctgctgc 60
actgccagcc tagggatgca cttgattccc aagaaatgca actgtcctat tcgcaragcc 120
gtccacaggt acctaccccc tggactgcag caactttatt accttaacta gcacaraaca 180
gaggttgatt taaactcctt acactcactt ctcaratcaa tgaatgggca aaraaacmcc 240
tcatggctct gggaaggcat gctgaracco gtttttgcaa gtccctgagga atggaaraat 300
atagctgcca ggtatcccaa gtctagggca gggagggkag tatcggcac ctttctactg 360
cattctgttg g
371

```

```

<210> 13
<211> 493
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(493)
<223> n = A,T,C or G

```

```

<400> 13
ccagtccaac ctgctcctca ttattgtata aatgagcaga atcaatatgg cggaagccag 60
ctycaattgc caatttgggt gcctctaaag ctttactttt aggaacctct gcaggcgcac 120
aggtgccaaa tcccaggaca ggcataaggt gaccatcatt cagcttcaca cactgatatt 180
tcgaatccat ttctgtcnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 240
caacctgctc ctcatatttg taaacatgtg cagaatcaat atggcggaac ccagcttcta 300
ttgctaattt tgtgacctcc aaagctttac ttctcggaac cttgggttctt ccgagcgctc 360

```


agcaatccccg ccgagcttct ttgagacgtc ctcaggtgtc ctttgacgat gcgtcctcca 420
 ctttcacaca ctctagcatt ccttcaactgg ggtcttcatt gccccacatt gggcagccag 480
 gaatgttggg gtg 493

<210> 14
 <211> 540
 <212> DNA
 <213> Homo sapien

<400> 14
 ccagatggtc cataatatgt caccgagcag gtgaatggca tttgtatgtc agccttggtt 60
 gtcttgact ccaggggtgga agtcatggta tagagctgag tcaactgggtc catttccttt 120
 ttaaaattat gaccaccgct ccttcaaggg gatgtagcac ttttccattc ctgtaccatg 180
 tgatattgcc atctggataa ctgtcttctg aaatgcagtc acccaacttt tttagctgct 240
 ctgtttcgag aaacagtgtc ttgcttaca tttcaggttt agatgggttg ttgaacacct 300
 tgactattgt aggtgcctca aacacgttgt cctcagttac tagcatgcac acaaactctc 360
 tttcatcact gatccttgca ttactgatag acaaagtgtg gttttctgag aggttcaatc 420
 tgtctttgta ttctggtaca tcgtcgtact gcacactttt cttttagtag gatctgaagg 480
 caataaatac tggggagcca tcgggctttt catattttcca tttgccc aaa catgagattc 540

<210> 15
 <211> 421
 <212> DNA
 <213> Homo sapien

<400> 15
 taccacctc cagcctccca tgtgagcctg tccttatgta tagtgtocaa cctctgattc 60
 tagcagtcaa gtgtcttccc caatcctaata gtccctgat atgtctctag cgaacttgacc 120
 atctcttggt ccttgggact ggggccagcc tcttgctgct ccaacttccct ctcatagtc 180
 agatagcccc aaaggctcta tcttttagctc ccagagaaact ttttggctct cagtatttcc 240
 cttccccctt ccttccattt ccccacaact gggggaggga agggagaaca ggggcacctg 300
 atcatcaatc tcccctgccc ctctcttgaa gccccctaga tttggatgaa gagcaggcca 360
 gtgagcaggg caaagcctgc taggagcaga atgaccttga ggatcctttg ctcagaactg 420
 g 421

<210> 16
 <211> 236
 <212> DNA
 <213> Homo sapien

<400> 16
 gccgtgtgtg cttttccag tgccgaggta cctatcgtc acggccagga gcttgctgtg 60
 gctgacagca aagagctgct ctctgtgggc ctgcttcac tcatccgaga ggccgtacaa 120
 gaagtgggtc attcctttgt ctgaaggagc gacaggagca tctacggttg agaagacaga 180
 aagtttggct tcgtcgtatg cttgctgtgt gaattttcca gacttagccc agtoga 236

<210> 17
 <211> 424
 <212> DNA
 <213> Homo sapien

<400> 17
 ccagaaaggt gacagtgggt ttccagggcc tcctgggcct ccaggtccac ctggtgaagt 60
 cattcagcct ttaccaatct tgtcctccaa aaaaacgaga agacatactg aaggcatgca 120

agcagatgca gatgataata ttcttgatta ctcgatgga atggaagaaa tatttggttc 180
 cctcaattcc ctgaaacaag acatcgagca tatgaaattt ccaatgggta ctcagaccaa 240
 tccagcccgga acttgtaaag acctgcaact cagccatcct gacttcccag atggtgaata 300
 ttggattgat cctaaccaag gttgctcagg agattccttc aaagtttact gtaatttcac 360
 atctggtggt gagacttgca tttatccaga caaaaaatct gagggagtaa gaatttcac 420
 atgg 424

<210> 18
 <211> 154
 <212> DNA
 <213> Homo sapien

<400> 18
 gtcaccaact ccttcagcgc ctccacaggg stttcggaca tgacagcaac cttttctccc 60
 aggacaattg aaatttgcta aagggaaagg ggaaagaaag ggaaaagga gaaaaagaa 120
 cacaagagac ttaaaggaca ggaggaggag atgg 154

<210> 19
 <211> 445
 <212> DNA
 <213> Homo sapien

<400> 19
 caacaaaatt ggtgaacaca tggaagaaca tggcatcaag tttataagac agttcgtacc 60
 aattaaagt gaacaaattg aagcagggac accaggccga ctcagagtag tagctcagtc 120
 caccaatagt gaggaatca ttgaaggaga atataatacg gtgatgctgg caataggaag 180
 agatgcttgc acaagaaaaa ttggcttaga aaccgtaggg gtgaagataa atgaaaagac 240
 tggaaaaata cctgtcacag atgaagaaca gaccaatgtg ccttacatct atgccattgg 300
 cgatatattg gaggataaagg tggagctcac ccagttgca atccaggcag gaagattgct 360
 ggctcagagg ctctatgcag gttccactgt caaagtgtga ctatgaaaat gttccaacca 420
 ctgtatttac tcctttggaa tatgg 445

<210> 20
 <211> 211
 <212> DNA
 <213> Homo sapien

<400> 20
 gggtgccact gctgcttga aagcactttc tgaacctaca gaagttgggt attgtctgaa 60
 atcccagagg acccataagt gccggtgaca agctgtctgt caggggagag gctccagaac 120
 ctgggttcgt cccagtgag accggaggat gatccccaa ggactgcgca gcatcagctc 180
 ttggtgggcc tctgccttct cttctgtttg g 211

<210> 21
 <211> 396
 <212> DNA
 <213> Homo sapien

<400> 21
 tgcccctgta ttggattgcc acacggctca cattgcatgc aagtttgctg agctgaagga 60
 aaagattgat cgccgttctg gtaaaaagct ggaagatggc cctaaattct tgaagtctgg 120
 tgatgctgcc attgttgata tggttcctgg caagcccatg tgtgttgaga gcttctcaga 180
 ctatccacct ttgggtcgct ttgctgttcg tgatatgaga cagacagttg cgggtgggtgt 240
 catcaaagca gtggacaaga aggctgctgg agctggcaag gtcaccaagt ctgccagaa 300

agctcagaag gctaaatgaa tattatccct aatacctgcc accccactct taatcagtgg 360
 tggaagaacg gtctcagaac tgtttgtttc aattgg 396

<210> 22
 <211> 277
 <212> DNA
 <213> Homo sapien

<400> 22
 ggaaccatgt ggccggcgcc cttgatcgtg agaaaggcga tgtgggagaa ctcccttcacg 60
 aagccggcaa tctgtctccc gctgtcccg tacttacta accagggcgg gcgctgcacc 120
 tccatcttct gggttgaggga atccacaaac cactcatccc ccatgaaatt gcaggccatg 180
 tctacatctc cattatataa taggatctgg gattttctgtg agctaagcag cttcagatac 240
 tgggagttca tgcttcggta gagacggcgg tactgta 277

<210> 23
 <211> 634
 <212> DNA
 <213> Homo sapien

<400> 23
 tctgaccatc catatccaat gttctcattt aaacattacc cagcatcatt gtttataatc 60
 agaaactctg gtccttctgt ctgggtggc ac tagagtctt ttgtgccata atgcagcagt 120
 atggagggag gattttatgg agaaatgggg atagtcttca tgaccacaaa taaataaagg 180
 aaaactaagc tgcatgtgtg gttttgaaaa gggttattata cttcttaaca attctttttt 240
 tcagggactt ttctagctgt atgactgtta cttgacctc tttgaaaagc attcccaaaa 300
 tgctctatct tagatagatt aacattaacc aacataatct tttttagatc gagtcagcat 360
 aaattttctaa gtcagcctct agtcgtgggt catctctttc acctgcattt tatttgggtg 420
 ttgtctgaag aaaggaaaga ggaaagcaaa tacgaattgt actattttgta ccaaactctt 480
 gggattcatt ggcaaataat ttcagtgtgg tgtattatta aatagaaaaa aaaaattttg 540
 tttcctaggt tgaaggtcta attgatacgt ttgacttatg atgaccattt atgcactttc 600
 aaatgaattt gctttcaaaa taaatgaaga gcag 634

<210> 24
 <211> 512
 <212> DNA
 <213> Homo sapien

<400> 24
 gcaaaacaag cctaagcaag cacaacgaag agcagaagtc agtgaaatta aaaagaggaa 60
 aaagaaaaat cataaaaaatc ataaaaagtt atttctttga aaagatcaat gaaatttagc 120
 aagactgaca cagataaaaa ggaatttagac ccaaactcagt gaacaggaat gaaatagagg 180
 atatcactac agaggctgca gccattgaaa ggataattag gaaatcccac agataacttt 240
 gtgtctcataa atttgacaat gtagaggaaa tatcttttagt ttttaattagc tttttatttt 300
 agttttttctc aaaaactaaa acttaataaaa actcaaccaa gacaaaatag acaatcagaa 360
 tgtaggcata cctcagagat gtggcggatt tggtttcaga ctactgcaat aaaccaata 420
 tggcaataaa aggagtcaca gaaagtgggt tcccagtgtg tatatataaa agttacattt 480
 actctatgaa gtgcaataac atttgtctaa aa 512

<210> 25
 <211> 461
 <212> DNA
 <213> Homo sapien

<400> 25

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctctgtttca | gcacctcatt | gggattattg | aactcattaa | attctttaca | tgaacttgaa | 60 |
| ttgttcattg | aaatctctag | ccatttccct | ggttaaacag | gataatcttt | ttttttcact | 120 |
| aaagaacatt | cgtgggtggt | tagtgatgag | gttaatatcc | ccctcttgtc | cacctccaca | 180 |
| ttggaaaaac | cacgttggac | tgagttttga | ggagcaaaga | actaatcact | tgaccaaagg | 240 |
| ggccctgtat | ccccacaagc | cctgggtatt | tttctctcat | agagagaaga | gggtctgtat | 300 |
| ggatacctga | aaatgtgatt | ttatatattc | ttggcatcca | ggggagaaaa | atcaaaaagc | 360 |
| aaggaagtta | cagttatctc | cccagaaatt | aatgggtcat | gtcaagacta | taggttttca | 420 |
| tttcttctg | ttgcttggtt | gaatgatggt | cttgtgggaa | a | | 461 |

<210> 26

<211> 317

<212> DNA

<213> Homo sapien

<400> 26

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tgctggagtc | ggaactgctg | cctttgtttg | goggccttgt | ttcttaaata | agttccctct | 60 |
| taggatttat | tacactaaaa | aaaaattagt | ttttgaaaag | aaataggaga | atacagaaac | 120 |
| atgaatttca | cgaggctatc | atctaacagt | gggggctttc | tacacacgtg | gtgccaaaat | 180 |
| gtgtcattct | gagtcaattg | caattcctct | ctaggagtga | aaagagataa | aagataagcc | 240 |
| aagaaccctg | gacagattct | tggtgttggt | gacaaagagg | aaaggacctg | agaatggggc | 300 |
| tggtggggag | aggggggg | | | | | 317 |

<210> 27

<211> 250

<212> DNA

<213> Homo sapien

<400> 27

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| taattgctgt | gattattaga | attctatcat | gactgtattg | tagtttttgc | tctattycag | 60 |
| ataagcmaga | tctaagaagt | tatcaaaact | attcttttaa | atgctaaagc | aggtaacttt | 120 |
| ttcttccatt | attttttcct | cctaccactg | agttttgtaa | tgaattcctt | gtgtatacaa | 180 |
| gcaatacagg | tgaataacta | actgttattt | ttagcttctt | caaaagctat | tttagaaaagc | 240 |
| ttcctggaaa | | | | | | 250 |

<210> 28

<211> 532

<212> DNA

<213> Homo sapien

<400> 28

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| cctatatcat | tcatttatac | agaagctgct | tgctgcttag | caagttggtg | ggtttgattt | 60 |
| tccttggttg | ctttgcagac | ctcccttgag | aggattcctt | ctggatggag | atttctttgt | 120 |
| tgctgtctcc | cttgccacaa | ctctgaccaa | gattgcattg | cgctatgtag | ctttggttca | 180 |
| ggagaagaaa | aagcaaaatt | cttttggtgc | tgaggctatg | ttgctcatgg | ctactatcct | 240 |
| gcatttggtg | aaatcctctc | ttcctaagaa | gcccaattact | gatgatgatg | tggatcgaat | 300 |
| ttccctgtgc | ctcaaggctc | tgtctgaatg | ttcaccttta | atgaatgaca | ttttcaataa | 360 |
| ggaatgcaga | cagtcccttt | ctcacatggt | atctgctaaa | ctagaagaag | agaaattatc | 420 |
| ccaaaagaaa | gaatctgaaa | agaggaatgt | gacagtacag | cctgatgacc | ccatttccct | 480 |
| catgcaacta | actgctaaga | atgaaatgaa | ctgcaaggaa | gatcagtttc | ag | 532 |

<210> 29

<211> 486

<212> DNA

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| ctgggcctcgg | atgggtctagg | atagccttac | tcacttgct | ggcaggtgac | aggctgttgg | 60 |
| ctggaattgc | ttggttctcc | tccatgtggc | ctctccagta | ggctagctca | ggcttattca | 120 |

<210> 38
 <211> 512
 <212> DNA
 <213> Homo sapien

<400> 38
 gctggaagtg aaatgcagat cagacccatt gtgatgtcac agaaagatgg ggacaggcca 60
 aagaaaaaag tgactttcaa ctcttcttcc atcattttta tcatcaccag tgatgaatca 120
 ctgtcagttg acgacagcga caaaaccaat ggggtccaaag ttgatgtaat ccaagttcgt 180
 cctttgtagg aatgaagaat ggcaacgaaa gatggggcct taaattggat gccacttttg 240
 gactttcatc ataagaagtg tctggaatac cgtttctatg taatatcaac agaaccttgt 300
 ggtccagcag gaaatccgaa ttgcccatat gctcttgggc ctccaggaaga ggttgaacaa 360
 aaacaaattc ttttaattca acgggtgctt tacataatga aaaaaccact tgtggcacac 420
 gatgggcac taaacatcatc atcttctaat gtgttggaga ttttcatttc aaatatattt 480
 tttaaattac tctattttcc aaaacacgta at 512

<210> 39
 <211> 370
 <212> DNA
 <213> Homo sapien

<400> 39
 ttttatgaac aagatataag gatcaaaaaa aagggtgttg atatgttttt ccaagcagag 60
 atgtactcga ctctgtccta tttagccttc ccataacctga cttctaatac cttttcctgg 120
 tgccttycca tctccctaac cccccctcac agggatgcct cctcccaagg ctccagaaac 180
 tctgacctc gcaactgttg agggagccca tgaattgctg gtcaatatcg ctcatcctct 240
 akactccatc ctgctgtgac ttcttcctac aagagctaga gaggcactga ctgataaata 300
 cctgtcacct gcccctttcc cagaggggtga aactccaccc actcccaactg cagaaatgaa 360
 tcttaaattg 370

<210> 40
 <211> 204
 <212> DNA
 <213> Homo sapien

<400> 40
 cctgaggggt ttccttttaa attttcattg agttgtccat ctccagcata tagggcttca 60
 ggagcagagc agacctgtt tttagtgggt ccatgggata aaatgggatt ggaggagcta 120
 gaagaattca gggctctggt caatctgcca gtcttcctga aatatcgaaa atacaccagg 180
 gctgctatat cagagccacc ctgg 204

<210> 41
 <211> 447
 <212> DNA
 <213> Homo sapien

<400> 41
 caggcagcaa ttcgtaaaga attaaatgag tacaaaagta atgaaatgga ggtacatgca 60
 tcaagcaagc acttgacaag attccacagg ccatagagat tttcttctga gaagaatttg 120
 tgtttaattt tttgatacca aactgaaca ttcacaggg aactttctctg aagttcagct 180
 caagactacc ctacctgtg tgtttgtgag aagagtagga tcacacacac aggtgcaatc 240
 ttgaccacac ttacctgcaa gaggagtaac cagaggacac acttccttcc ttctttggtg 300
 tctgaggagt gtgaactgtt ggggtcagtt aagacccaac ataactctat cagaagaaaa 360

ctgttgtttg cttttcaacc ttgttttaca gttctgcagt gtagtggagg acgggcaacg 420
 tgcattgtgca ggctcaccac tcccagg 447

<210> 42
 <211> 498
 <212> DNA
 <213> Homo sapien

<400> 42
 ctggttttgt aaaaacagtc tctttattct actgtgctga aacctcacc aatatagaaa 60
 attagattct cattgcactg aactatattt atatgcctaa gtatgtagaa gtaaaattat 120
 ataccccaaa aggattttat cttgttgtat atattaaatg ttatttctgc atatagggtc 180
 ttttatggag aaactgatga tgataagctt aatactcact tgtttagcag catctgaatg 240
 cacaaatgct ttatatatct cttctgcttt acagggcaaa agatcagact ctgttttctt 300
 atagtcttca caagccagcc agaactcaat attctcctca ctgaattcag actttaggaa 360
 acttccaaag acattttgac cagtttgggt ggcaagaagt tttccagag attgagacca 420
 ttgcattact tcagcagcag aaagtacatc cttggacttg gaagatttca ttccagattc 480
 cagatgtggg atcataga 498

<210> 43
 <211> 312
 <212> DNA
 <213> Homo sapien

<400> 43
 caggaaggcg gccagaatg tgagtgcaaa gattggttcc tgagagcccc gagaagaaaa 60
 ttcattgacag tgtctgggct gccaaagaag cagtgcacct gtgatcattt caagggcaat 120
 gtgaagaaaa caagacacca aaggcaccac agaaagccaa acaagcattc cagagcctgc 180
 cagcaatttc tcaaacaatg tcagctaaga agctttgtct tgcccttgta ggagctctga 240
 gcgcccactc ttccaattaa acattctcag ccaagaagac agtgagcaca cctaccagac 300
 actctttctt tc 312

<210> 44
 <211> 417
 <212> DNA
 <213> Homo sapien

<400> 44
 ctaacacatt tactctccac tattcgtact ctggtagcca tgtaacccc atcagagatt 60
 ccttctcaag ccatgtctca gagctgagag gcatcccagc aagttttgca gctcacagtt 120
 ttttccgtaa attacttatt ctataaaatt ggagtaggcc ataaactttg gagggcccta 180
 gaccaatttt ttggattatt tttcgtcttc tatcattccg ctgatcttag atattctctg 240
 cattaaatat taaatatcac ttctaggttg aaaaatcccc ctaaaaatat ttctagctca 300
 gatttttctt ccaaattctg caatagaaga tcacaatgtg aactctgcat ctccatgtta 360
 aagtctaattg gacattcaca cttagcatgt ctcaaagaaa tctcatgtaa accatgg 417

<210> 45
 <211> 494
 <212> DNA
 <213> Homo sapien

<400> 45
 cgcggtgtctg tgggtatgtg acacgtgcat gttctgcatg tctgtaggtc acacatgctt 60
 tgggtgcatgt acacgtgtgt gtgtgtatgc gtgtaggagc tcacacttgt gtacacgttt 120


```

gtgtgcatgc atgtgtgcag gagcttgca c gtttgtggtg ggtacatgta catatgtgag 180
tgatcctgtg tgcaagcccc catgtggaca tggctatgag tgagcgtgga gccaaaagcc 240
aggtaacacg catgcagcag gccactgtg cgtgtctgag acggtctgtg gcagggactg 300
ggtgtgaatc atgcagcagg cccactgtgc gtgtctgaga cggctctgtg cagggactgg 360
gtgtgaatca gtgaccgtgt ctctgaccaa catgctgaat tacaaattga taatttatta 420
acctgtgcag caacaaataa gatttttcaa aactcaacaa agtgctcaaa gttgacatta 480
cttgcttcaa agtt 494

```

```

<210> 46
<211> 516
<212> DNA
<213> Homo sapien

```

```

<400> 46
ccagtccaac ctgctcctca ttattgtata aatgagcaga atctatatgg cggaacccag 60
cttctattgc taattttgtg acctocaaag ctttacttct cggaacctcc tcctttggcc 120
gtcatttgat cattcaactc tttgtcagtg gcaactcccg ctattttggt gtgttggttt 180
gttactacac agtgagcaca aacatggtgg tccaatacag aggcctcttc tgtcaggtgt 240
caaccagaaa gttcatctaa cactgtgata tttgcatcct tcttgaacag ttgttggtctg 300
aagattcatt tgatgaatcg atttttcaaa agagatgatt cttggttctt ccgagcgctc 360
agctctcccg ccgagcttct ttgagacgtc ctcaggtgtc ctttgacgat gcgtcctcca 420
ctttcacaca ctctagcatt ccttacttgg ggtcttcatt gcccacatt gggcagccag 480
gaatgttggg gtgatcagac acaacaccag gtcatg 516

```

```

<210> 47
<211> 459
<212> DNA
<213> Homo sapien

```

```

<400> 47
ccaattcaga gtggcattct gcatttctgt ggcttccaag tcttagaacc tcaactgaca 60
tatagcattg ggcacactcc agcagacgcc cgaattcaaa tcttggaagg atggaagaaa 120
cgcttgaga atatttgga tgagacacca ctgtattttg ctccaagcag cctctttgac 180
ctaaacttcc aggcaggatt cttaatgaaa aaagaggtac aggatgagga gaaaaacaag 240
aaatttgccc tttctgtggg ccatacattg ggcaagtcca tcccaactga caaccagatc 300
aaagctagaa aatgagattc cttagcctgg atttcttct aacatgttat caaatctggg 360
tatctttcca ggcttccctg acttgcttta gtttttaaga tttgtgtttt tctttttcca 420
caaggaataa atgagaggga atcgaksaaa aaaaaaaaa 459

```

```

<210> 48
<211> 430
<212> DNA
<213> Homo sapien

```

```

<400> 48
cctatatcca gccacagcct ctgggagtgg tgctgataat cggagcttgg aattaccctt 60
tcgtttcac cattcagcca ctgataggag ccatcgctgc aggaaatgct gtgattataa 120
agccttctga actgagtga aatacagcca agatcttggc aaagcttctc cctcagtatt 180
tagaccagga tctctatatt gttattaatg gtggtgttga ggaaaccacg gagctcctga 240
agcagcgatt tgaccacatt ttctatacgg gaaacactgc ggttggcaaa attgtcatgg 300
aagctgctgc caagcatctg acccctgtga ctcttgaact gggagggaaa agtccatgtt 360
atattgataa agattgtgac ctggacattg tttgcagacg cataacctgg ggaaaataca 420
tgaattgtgg 430

```

<210> 49
 <211> 288
 <212> DNA
 <213> Homo sapien

<400> 49
 ccatccgaag caagattkca gatggcagtg tgaagagaga agacatattc tacacttcaa 60
 agctttggwg caattcccat cgaccagagt tgggccgacc agccttggaa aggtcactga 120
 aaaatcttca attggattat gttgacctct accttattca ttttccagtg tctgtaaagc 180
 caggtgagga agtgatccca aaagatgaaa atggaaaaat actatttgac acagtggatc 240
 tctgtgccac gtgggaggcc rtggagaagt gtaaagatgc aggattgg 288

<210> 50
 <211> 411
 <212> DNA
 <213> Homo sapien

<400> 50
 ccagagaatg acattcatgt ccccgaggat cccttgacaga gaggatcatg agccactgcc 60
 accagtggtg atggaaagca ctgtcttctt actccggaag ggctccttctg catacatggc 120
 agcgtaagt taagcaaact ctctatgaa cactcgtcga aaccagcctt tcagaatggc 180
 agggactcca aaccactgca gggggaactg gaatatcaca aggtctgcgg cttccagctt 240
 cttttgttca gccacaatat ctgggctcag atggccttct ttataagcca gaacagactc 300
 ggcaggatac tgaaagttcg cagggtcctt cagtttacct gtgatgtcct ttctggaaat 360
 gatgggattg aagttcatgg catagaggtc cgactccacc acctcccatc c 411

<210> 51
 <211> 503
 <212> DNA
 <213> Homo sapien

<400> 51
 gatattcttat gattaataaac aaattaaatt ttaaaacacc tgaagatata ttagaagaaa 60
 ttgtgcaccc tccacaaaac atacaaagtt taaaagtttg gatctttttc tcagcaggta 120
 tcagttgtaa ataataaatt aggggccaaa atgcaaaacg aaaaatgaag cagctacatg 180
 tagttagtaa tttctagttt gaactgtaat tgaatattgt ggcttcatat gtattatattt 240
 atattgtact tttttcatta ttgatggttt ggactttaat aagagaaatt ccatagtttt 300
 taatatccca gaagtgcagc aatttgaaca gtgtattcta gaaaacaata cactaactga 360
 acagaagtga atgcttatat atattatgat agccttaaac ctttttcctc taatgcctta 420
 actgtcaaat aattataacc ttttaaagca taggactata gtcagcatgc tagactgaga 480
 ggtaaactat gatgcaatta aga 503

<210> 52
 <211> 503
 <212> DNA
 <213> Homo sapien

<400> 52
 gatattcttat gattaataaac aaattaaatt ttaaaacacc tgaagatata ttagaagaaa 60
 ttgtgcaccc tccacaaaac atacaaagtt taaaagtttg gatctttttc tcagcaggta 120
 tcagttgtaa ataataaatt aggggccaaa atgcaaaacg aaaaatgaag cagctacatg 180
 tagttagtaa tttctagttt gaactgtaat tgaatattgt ggcttcatat gtattatattt 240
 atattgtact tttttcatta ttgatggttt ggactttaat aagagaaatt ccatagtttt 300
 taatatccca gaagtgcagc aatttgaaca gtgtattcta gaaaacaata cactaactga 360

acagaagtga atgcttatat atattatgat agccttaaac ctttttcctc taatgcctta 420
 actgtcaaat aattataacc ttttaaagca taggactata gtcagcatgc tagactgaga 480
 ggtaaact gatgcaatta aga 503

<210> 53
 <211> 531
 <212> DNA
 <213> Homo sapien

<400> 53
 tttttttttt tttttaaaat gaggatattt tattatttca ggtaattttc ccagaggkga 60
 gaatagtaca tgggaaattc tctttaggcc aggtctagta ttacagkgtg gkgctcaagg 120
 ccgcccatac gaacagtgat actctcccaa cagatttcat ccaccccgtc tccactaact 180
 tttgccataa aaattcctct gaattgtatc ttcttggaag aagtaaataat ctgttcgact 240
 atacaaagaa acagagaaac cactcccatt gcaatcaatc ttcaagagag ggagcaggca 300
 agccgtgttc tttctgctga gttttataga ctctgacaag ctgtgaaata aacataaaca 360
 gaagacaaaa cagtgcacaa aataagcagt agatgaccct gtgacaagac ggcattgcag 420
 aacaaagact gacgttttaa ggggagtcac gcagagtaac atgggaacac aagcctgaca 480
 acctggtcag cttccactta ctctagctcc tttgaactct caacactaaa a 531

<210> 54
 <211> 450
 <212> DNA
 <213> Homo sapien

<400> 54
 ccatgggtgt ctggagcwcc ctgaaactgt atcaaagttg tacatatttc caaacatttt 60
 taaaatgaaa aggcactctc gtgttctcct cactctgtgc actttgctgt tgggtgtgaca 120
 aggcatttaa agatgtttct ggcattttct ttttatttgt aagggtggtg taactatggt 180
 tattggctag aaatcctgag ttttcaactg tatatatcta tagtttgtaa aaagaacaaa 240
 acaaccgaga caaacccttg atgctccttg ctggcgcttg aggcctgtggg gaagatgcct 300
 tttgggagag gctgtagctc agggcggtgc ctgtgaggct ggacctgttg actctgcagg 360
 gggcatccat ttagcttcag gttgtcttgt ttctgtatat agtgacatag cattctgctg 420
 ccatcttagc tgtggacaaa ggggggtcag 450

<210> 55
 <211> 648
 <212> DNA
 <213> Homo sapien

<400> 55
 caacttcaac cacaggctgc tggasatgat cctcarcaag ccagggtcga agtacaagcc 60
 tgtctgcaac caggtggaat gtcatectta cttcaaccag agaaaaactgc tggatttctg 120
 caagtcaaaa gacattgttc tggttgcta tagtgctctg ggatcccacc gagaagaacc 180
 atgggtggac ccgaactccc cgggtgctctt ggaggaccca gtcctttgtg ccttggcaaa 240
 aaagcacaag cgaacccag cctgattgc cctgcgctac cagctrcagc gtgggggtgt 300
 ggtcctggcc aagagctaca atgagcagc catcagacag aacgtgcagg tgtttgaatt 360
 ccagttgact tcagaggaga tgaaagccat agatggccta aacagaaatg tgcgatattt 420
 gacccttgat atttttgctg gccccctaa ttatccattt tctgatgaat attaacatgg 480
 agggcattgc atgaggtctg ccagaaggcc ctgcgtgtgg atggtgacac agaggatggc 540
 tctatgctgg tgaactggaca catcgctctt ggtaaactct ctctgcttg gygayttcag 600
 caagctacag caaagcccat tggccggaat aaatatcaag ggtcaaat 648

<210> 56

<211> 536
 <212> DNA
 <213> Homo sapien

<400> 56
 ctggcatgag aatatttttt tttttaagtg cggtagtttt taaactgttt gtttttaaac 60
 aaactataga actcttcatt gtcagcaaag caaagagtca ctgcatcaat gaaagttcaa 120
 gaacctcctg tacttaaaca cgattcgcaa cgttctgtta ttttttttgt atgttttagaa 180
 tgctgaaatg tttttgaagt taaataaaca gtattacatt tttaaaactc ttctctatta 240
 taacagtcaa tttctgactc acagcagtga acaaaccccc actccattgt atttggagac 300
 tggcctccct ataaatgtgg tagcttcttt tattactcag tggacctgcc cgggcggccg 360
 ctggaagccg aattccagca cactggcggc cgttactagt ggatccgagc tcggtaccaa 420
 gcttggccgt aatcatggtc atagctgttt cctgtgtgaa attgttatcc gtcacaatt 480
 ccacacaaca tacgagccgg aagcataaag tgtaaagcct ggggtgccta atgagt 536

<210> 57
 <211> 391
 <212> DNA
 <213> Homo sapien

<400> 57
 aggaactact gtcccagagc tgaggcaagg ggattttctca ggtcatttgg agaacaagtg 60
 ctttagtagt agtttaaagt agtaactgct actgtattta gtggggtgga attcagaaga 120
 aatttgaaga ccagatcatg ggtggtctgc atgtgaatga acaggaatga gccggacagc 180
 ctggctgtca ttgctttctt cctccccatt tggacccttc tctgccctta catttttgtt 240
 tctccatcta ccaccatcca ccagtctatt tatttgtcta gttggatttc atttctcttg 300
 gaaaatttat tgtttattgg catgtgacct ttgactgatg gcttcattag cattytgttt 360
 ttcttttttg atccttaata gaaaactcaa t 391

<210> 58
 <211> 455
 <212> DNA
 <213> Homo sapien

<400> 58
 gaagacatgc ttacttcccc ttcaccttcc ttcattgatg gggaagagtg ctgcaaccca 60
 gccctagcca acgccgcatg agagggagtg tgccgagggc ttctgagaag gtttctctca 120
 catctagaaa gaagcgctta agatgtggca gccctcttc ttcaagtggc tcttgtcctg 180
 ttgccctggg agttctcaaa ttgctgcagc agcctccacc cagcctgagg atgacatcaa 240
 tacacagagg aagaagagtc aggaaaagat gagagaagtt acagactctc ctgggcgacc 300
 ccgagagctt accattcctc agacttcttc acatggtgct aacagatttg ttcctaaaag 360
 taaagctcta gaggccgtca aattggcaat agaagccggg ttccaccata ttgattctgc 420
 acatgtttac aataatgagg agcaggttgg actgg 455

<210> 59
 <211> 398
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(398)
 <223> n = A,T,C or G

<400> 59

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|-----|
| ctcagaggca | gcggtgcgggt | gtgctctttg | tgaaattcca | ccatggcgta | ccgtggccag | 60 |
| ggtcagaaag | tgcagaaggt | tatgggtgcag | cccatcaacc | tcctcttcag | atacttacia | 120 |
| aatagatcgc | ggattcaggt | gtggctctat | gagcaagtga | atatgcggat | agaaggctgt | 180 |
| atcattgggt | ttgatgagta | tatgaacctt | gtattagatg | atgcagaaga | gattcattct | 240 |
| aaaacaaagt | caagaaaaca | actngntcgg | atcatgctaa | aaggagataa | tattactctg | 300 |
| ctacaaagtg | tctccaacta | gaaatgatca | atgaagtga | aaattggtga | gaaggataca | 360 |
| gtttgttttt | agatgtcctt | tgtccaatgt | gaacattt | | | 398 |

<210> 60

<211> 532

<212> DNA

<213> Homo sapien

<400> 60

| | | | | | | |
|------------|------------|-------------|-------------|------------|------------|-----|
| gacttctgag | acctggggca | cccgggcctt | tgcggcagct | actggcaggg | cctggccacc | 60 |
| tcataggact | cagttccctt | ctgaacactc | gggggacatg | ggcctctaac | tgccactct | 120 |
| gatatgcctg | ggtgagccta | ggaggggaagg | ctctgatattg | gatttctcca | gtcaaagctc | 180 |
| acagaaaaaa | acctggcact | ttgattttca | tgggatggtc | ctaacagggg | cagtcacctc | 240 |
| cgagcagttt | gggaacccag | tttcttgtcc | tgggcccctca | ggtcagcctg | gctgaattag | 300 |
| gaccttccct | tggcacaggg | gtgagaaaga | gcttggggaa | cgcttggcat | tatggagggc | 360 |
| tgggaagggc | tcaaccccga | tttggagaga | agtttgggat | ggagtgggcg | agagattgag | 420 |
| agagcgagca | ggaaaagagg | tcttggagcc | tgggactgat | ggtggataag | gcctggaaag | 480 |
| aasatgacsa | ggaggaggag | agaggggaagt | gggtggatga | ggagcaggct | ga | 532 |

<210> 61

<211> 466

<212> DNA

<213> Homo sapien

<400> 61

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| gcgacggcga | cgtctctttt | gactaaaaga | cagtgtccag | tgctccagcc | taggagtcta | 60 |
| cggggaccgc | ctcccgcgcc | gccacatgc | ccaacttctc | tggcaactgg | aaaatcatcc | 120 |
| gatcggaana | cttcgaggaa | ttgctcaaa | tgctgggggt | gaatgtgatg | ctgaggaaga | 180 |
| ttgctgtggc | tgcagcgtcc | aagccagcag | tggagatcaa | acaggagggg | gacactttct | 240 |
| acatcaaaac | ctccaccacc | gtgcgcacca | cagagattaa | cttcaagggt | ggggaggagt | 300 |
| ttgaggagca | gactgtggat | gggagggcct | gtaagagcct | ggtgaaatgg | gagagtgaga | 360 |
| ataaaatgg | ctgtgagcag | aagctcctga | agggagaggg | ccccagacc | togtggacca | 420 |
| gagaactgac | caacgatggg | gaactgatcc | tgaccatgac | ggcgga | | 466 |

<210> 62

<211> 548

<212> DNA

<213> Homo sapien

<400> 62

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ttttgaattt | acaccaagaa | cttctcaata | aaagaaaatc | atgaatgctc | cacaatttca | 60 |
| acataaccaca | agagaagtta | atttcttaac | attgtgttct | atgattattt | gtaagacctt | 120 |
| caccaagttc | tgatatcttt | taaagacata | gttcaaaatt | gcttttgaaa | atctgtattc | 180 |
| ttgaaaatat | ccttggttgt | tattaggttt | ttaaatacca | gctaaaggat | tacctcactg | 240 |
| agtcacagct | accctcctat | tcagctcccc | aagatgatgt | gtttttgctt | accctaagag | 300 |
| aggttttctt | cttattttta | gataattcaa | gtgcttagat | aaattatgtt | ttctttaagt | 360 |
| gtttatggta | aactctttta | aagaaaattt | aatatgttat | agctgaatct | ttttggtaac | 420 |
| tttaaattct | tatcatagac | tctgtacata | tgttcaaatt | agctgcttgc | ctgatgtgtg | 480 |


```
<210> 66
<211> 535
<212> DNA
<213> Homo sapien
```

```
<210> 67
<211> 527
<212> DNA
<213> Homo sapien
```

```
<210> 68
<211> 431
<212> DNA
<213> Homo sapien
```

| <400> 68 | | | | | | |
|-------------|-------------|-------------|-------------|------------|------------|-----|
| gggaaacttc | atgggtttcc | tcattctgtca | tgtcgatgat | tatatatgga | tacatttaca | 60 |
| aaaataaaaa | gcgggaattt | tcccttcgct | tgaatattat | cctgttatat | tgcatgaatg | 120 |
| agagatttcc | catatttcca | tcagagtaat | aaatataact | gctttaattc | ttaagcataa | 180 |
| gtaaacaatga | tataaaaata | tatgctgaat | tacttgtgaa | gaatgcattt | aaagctattt | 240 |
| taaatgtgtt | tttatttgta | agacattact | tattaagaaa | ttggttatta | tgcttactgt | 300 |
| tctaactctgg | tggtaaagggt | attcttaaga | atttgcaggt | actacagatt | ttcaaaactg | 360 |
| aatgagagaa | aattgtataa | ccatcctgct | gwtccttttag | tgcaatacaa | taaaactctg | 420 |

aaattaaaac t

431

<210> 69

<211> 399

<212> DNA

<213> Homo sapien

<400> 69

| | | | | | | |
|------------|------------|------------|-------------|-------------|-------------|-----|
| gacacggcgg | acacacacaa | acacagaacc | acacagccag | tcccaggagc | ccagtaatgg | 60 |
| agagcccaa | aaagaagaac | cagcagctga | aagtccggat | cctacacctg | ggcagcagac | 120 |
| agaagaagat | caggatacag | ctgagatccc | agtgcgcgac | atggaagggtg | atctgcaaga | 180 |
| gctgcatcag | tcaaacaccg | gggataaatc | tggattttggg | ttccggcgtc | aagggtgaaga | 240 |
| taatacctaa | agaggaacac | tgtaaaatgc | cagaagcagg | tgaagagcaa | ccacaagttt | 300 |
| aatgaagac | aagctgaaac | aacgcaagct | ggttttatat | tagatatattg | acttaaacta | 360 |
| tctcaataaa | gttttgcagc | tttcaccaar | aaaaaaaaa | | | 399 |

<210> 70

<211> 479

<212> DNA

<213> Homo sapien

<400> 70

| | | | | | | |
|-------------|------------|------------|-------------|------------|-------------|-----|
| cgcggcggag | ctgtgagccg | gcgactcggg | tccctgaggt | ctggattctt | tctccgctac | 60 |
| tgagacacgg | cggacacaca | caaacacaga | accacacagc | cagtcccagg | agcccagtaa | 120 |
| tggagagccc | caaaaagaag | aaccagcagc | tgaaagtccg | gatacctaac | ctgggcagca | 180 |
| gacagaagaa | gatcaggata | cagctgagat | cccagggtgct | gggaagggaa | atgcgcgaca | 240 |
| tggaagggtga | tctgcaagag | ctgcatcagt | caaacacccg | ggataaatct | ggattttgggt | 300 |
| tccggcgctca | aggtgaagat | aatacctaaa | gaggaacact | gtaaaatgcc | agaagcaggt | 360 |
| gaagagcaac | cacaagttta | aatgaagaca | agctgaaaca | acgcaagctg | gttttatatt | 420 |
| aggatatattg | acttaaacta | tctcaataaa | gttttgcagc | tttcaccaa | aaaaaaaaa | 479 |

<210> 71

<211> 437

<212> DNA

<213> Homo sapien

<400> 71

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ctcagcggct | gccaacagat | catgagccat | cagctcctct | ggggccagct | ataggacaac | 60 |
| agaactctca | ccaaaggacc | agacacagtg | rgcaccatgg | gacagtgtcg | gtcagccaac | 120 |
| gcagaggatg | ctcaggaatt | cagtgatgtg | gagagggcca | ttgagaccct | catcaagaac | 180 |
| tttcaccagt | actccgtgga | gggtgggaag | gagacgctga | ccccttctga | gctacgggac | 240 |
| ctggtcaccc | agcagctgcc | ccatctcatg | ccgagcaact | gtggcctgga | agagaaaatt | 300 |
| gccaacctgg | gcagctgcaa | tgactctaaa | ctggagttca | ggagtttctg | ggagctgatt | 360 |
| ggagaagcgg | ccaagagtgt | gaagctggag | aggcctgtcc | gggggcaactg | agaactccct | 420 |
| ctggaattct | tggggggg | | | | | 437 |

<210> 72

<211> 561

<212> DNA

<213> Homo sapien

<400> 72

| | | | | | | |
|-------------|------------|------------|-------------|------------|------------|-----|
| ggatgggtata | ctgtaaattc | agcatatgga | gataaccatta | tcataccttg | ccgacttgac | 60 |
| gtacctcaga | atctcatgtt | tggcaaatgg | aaatatgaaa | agcccgatgg | ctccccagta | 120 |


```
<210> 73
<211> 916
<212> DNA
<213> Homo sapien
```

```
<210> 74
<211> 547
<212> DNA
<213> Homo sapien
```

```
<210> 75
<211> 793
<212> DNA
<213> Homo sapien
```

<400> 75

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| tgaggaagtt | gcaagccaac | aaaaaagttc | aaggatctag | aagacgatta | aggggaaggtc | 60 |
| gttctcagt | aaaatccaaa | aaccagaaaa | aatgtttat | acaaccctaa | gtcaataacc | 120 |
| tgaccttaga | aaattgtgag | agccaagttg | acttcaggaa | ctgaaacatc | agcaciaaaga | 180 |
| agcaatcatc | aaataattct | gaacacaaat | ttaatatatt | tttttctgaa | tgagaaacat | 240 |
| gagggaaatt | gtggagttag | cctcctgtgg | agttagcctc | ctgtggtaaa | ggaattgaag | 300 |
| aaaatataac | accttacacc | ctttttcatc | ttgacattaa | aagttctggc | taactttgga | 360 |
| atccattaga | gaaaaatcct | tgtcaccaga | ttcattacaa | ttcaaactga | agagttgtga | 420 |
| actgttatcc | cattgaaaag | accgagcctt | gtatgtatgt | tatggatata | taaaatgcac | 480 |
| gcaagccatt | atctctccat | gggaagctaa | gttataaaaa | taggtgcttg | gtgtacaaaa | 540 |
| ctttttatat | caaaaaggctt | tgacacattc | tatatgagtg | ggtttactgg | taaattatgt | 600 |
| tattttttac | aactaatatt | gtactctcag | aatgtttgtc | atatgcttct | tgcaatgcat | 660 |
| attttttaat | ctcaaacggt | tcaataaaac | catttttcag | atataaagag | aattacttca | 720 |
| rattgagtaa | ttcagaaaaa | ctcaagattt | aagttaaaaa | gtgggttgga | cttggaaca | 780 |
| ggactttata | cct | | | | | 793 |

<210> 76

<211> 461

<212> DNA

<213> Homo sapien

<400> 76

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| accttgact | attcccctca | gtccatctat | cgaggtcttt | gcaggaagca | tactgggaat | 60 |
| tgaaacgaga | gcctaaatga | catctaagaa | aggcagtggt | caataaccagg | tattaggtga | 120 |
| ggatgggatt | ctaaggacat | cagtgggagg | cagggagcca | ccttcagacc | tcagcatgga | 180 |
| agcttccaag | atccagagga | agaggcaaca | gcactgagag | tcataggtag | aagaatcatc | 240 |
| acagccctgc | taaccaggca | gctgatgccc | ctctcccctg | gctccctgtg | tccaaatcct | 300 |
| acaggggcat | ctgttggctg | aactcaacct | gaagccaaag | agaagatgag | tggagagagg | 360 |
| caacatttat | agagctcagg | tttctagggc | tggagaggga | tctggaggga | cacacaggag | 420 |
| acacctggca | taaccaaaaa | atgattaaaa | aaaaaaaaaa | a | | 461 |

<210> 77

<211> 642

<212> DNA

<213> Homo sapien

<400> 77

| | | | | | | |
|-------------|------------|------------|-------------|------------|-------------|-----|
| ggttgacga | aacacactgg | ggaatggagc | aaaacagtct | ttgaatatcg | aacacgcaag | 60 |
| gctgtgagac | tacctattgt | agatattgca | ccctatgaca | ttggtggtcc | tgatcaagaa | 120 |
| tttgggtggtg | acgttggccc | tgtttgcttt | ttataaaacca | aactctatct | gaaatcccaa | 180 |
| caaaaaaaat | ttaactccat | atgtgttcct | cttgttctaa | tcttgtcaac | cagtgtcaagt | 240 |
| gaccgacaaa | attccagtta | tttatttcca | aatgttttgg | aaacagtata | atttgacaaa | 300 |
| gaaaaatgat | acttctcttt | ttttgctgtt | ccaccaata | caattcaa | gctttttgtt | 360 |
| ttattttttt | accaattcca | atttcaaaat | gtctcaatgg | tgctataata | aataaacttc | 420 |
| aacactcttt | atgataacaa | aaaaaarawa | wattctttga | atcctagccc | atctgcagag | 480 |
| caatgactgt | gctcaccagt | aaaagataac | ctttctttct | gaaatagtca | aatacgaaat | 540 |
| tagaaaagcc | ctccctattt | taactacctc | aactggtcag | aaacacagat | tgtattctat | 600 |
| gagtcccaga | agatgaaaaa | aattttatac | gttgataaaa | ct | | 642 |

<210> 78

<211> 519

<212> DNA

<213> Homo sapien

<400> 78
gcagaagaag aagcggacct tccgcaagtt cacctaccgc ggcgtggacc tcgaccagct 60
gctggacatg tcctacgagc agctgatgca gctgtacagt gcgcgccagc ggcggcggct 120
gaaccggggc ctgcggcgga agcagcactc cctgctgaag cgcttgcgca aggccaagaa 180
ggaggcgccg cccatggaga agccggaagt ggtgaagacg cacctgcggg acatgatcat 240
cctacccgag atggtgggca gcatggtggg cgtctacaac ggcaagacct tcaaccaggt 300
ggagatcaag cccgagatga tcggccacta cctgggcgag ttctccatca cctacaagcc 360
cgtaaagcat ggccggcccg gcatcggggc caccactcc tcccgttca tccctctcaa 420
gtaatggctc agctaataaa aggcgcacat gactccaaaa aaaaaaaaaa aaggcgggcc 480
gccaccgcgg gggagctcca cttttgttcc ctttaatga 519

<210> 79
<211> 526
<212> DNA
<213> Homo sapien

<400> 79
gtctggaggc ggtgtcctct ccgccctgtc gggctcctgga tgagtacgag ttatgggtcac 60
ggtcacagcc tgatctctta tgtgttcata gccattcgct ctcccatcag aactgtttgt 120
cctgaatgtg ttctctagt tctagaaaat gaccactaat ttaaaaaact cggttgtgag 180
gtttgccagc aggcacttgt tccagaattt cccctcctgc ttcagccatg tccttgtcac 240
ttggcattct aagctaaaagc tttagcttcc caattogtga tgtgctaggc caagattcgg 300
gagctgttgc cagcctcgtc aaatatggaa gagaaacaac ctgcgggtcaa aaggagtgta 360
tttgtttaagt ggtgcgcgtc tatctcataa ctagatgtac caaccaggga agggccaagg 420
atggaaaggg gtaacttttg tgcttccaaa gtagctaagc agaagtgggg gagcagttta 480
gccagatgat ctttgattag gcaaacattg agttttaaag aggctg 526

<210> 80
<211> 281
<212> DNA
<213> Homo sapien

<400> 80
gttatattag tgggtagtgt aacattttat ccaggttggg gtgaggggag atggccacag 60
tagcaagtgg tgacactaaa taccattttg aaggctgatg tgtatataca tcattactgt 120
ccgtagcaat gaaggataca gtactgtgtt gtgggtgagt gttgctattg cccagcatta 180
atatttgggt gtgtatgttt gaggctatga aacacgcagg agtggttttt tgctattaat 240
tttaagagaa agcagctttt tcttaaaatt cactgttgag a 281

<210> 81
<211> 405
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(405)
<223> n = A,T,C or G

<400> 81
gtgggtggga gcgcgtgctg ttgggagttg cttggagggt ggccggcgcg ggctgaaggc 60
tagcaaaccg agcgatcatg tcgcacaaac aaatttacta ttccgacaaa tacgacsacg 120
aggagtttga statcgacat gtcattgctg ccaaggacat akccaasctg gtccctaaaa 180
cccatctgat gtctgaatct gaatggagga atcttggcng ttcagmagan tcagggatgg 240

<211> 401
 <212> DNA
 <213> Homo sapien

<400> 85
 cagtgtggtg gaattcccaa gatagaaatg aaaaactctt ttatagagtg ctgacatctg 60
 acattgagaa attcatgcct attgtttata ctcccactgt gggctctggct tgccaacaat 120
 atagtttggt gtttcggaag ccaagagggtc tctttattac tatccacgat cgagggcata 180
 ttgcttcagt tctcaatgca tggccagaag atgtcatcaa ggccattgtg gtgactgatg 240
 gagagcgtat tcttggttg ggagaccttg gctgtaatgg aatgggcatc cctgtgggta 300
 aattggctct atatacagct tgcggaggga tgaatcctca agaatgtctg cctgtcatte 360
 tggatgtggg aaccgaaaat gaggagttac ttaaagatcc a 401

<210> 86
 <211> 547
 <212> DNA
 <213> Homo sapien

<400> 86
 gaagcctctt gtgtttgtgt gcagagaagt atatgatcca ccatgctaata gacacttgcc 60
 tttttttcca ccattaaggc tttaagaaca tgtggaataa gtttttttagc tgctaatagac 120
 aaaacaaatc ctgtaactac ccagccagca agtatatagc acagaacact gtgttacttt 180
 acaagggtct atgtgactgg aataaggtgg tcccacttga ctgttccaaa gagcagcttc 240
 tcagatcttc agtgttcact ggtaaatttc taacagtgtg tttgtgtaaa gtttgtcatt 300
 tcatactcca tacactacag ttgctgtcac tgatccctgt tttgctggct ttttaagctac 360
 ttggtcaaaa atcctgcttc cttaaaacat agagaattaa tgagcatctc aagctttttc 420
 ttttcctttt taatgatgac tgcactatca agagtattct agtgttctct ctttgttttg 480
 catataatca tgcaccaaac tttttatttc ttttaaggtgg gagtatattt ttatttccta 540
 aatgcca 547

<210> 87
 <211> 530
 <212> DNA
 <213> Homo sapien

<400> 87
 atggattcga aataccagkg tgtgaagctg aatgatggtc acttcatgcc tgtcctggga 60
 tttggcacct atgcgcctgc agaggttcct aaaagtaaaag ctctagaggc cgtcaaattg 120
 gcaatagaag ccgggttcca ccatattgat tctgcacatg tttacaataa tgaggagcag 180
 gttggactgg ccatccgaag caagattgca gatggcagtg tgaagagaga agacatattc 240
 tacacttcaa agcttttgag caattcccat cgaccagagt tggtcgacc agccttgga 300
 aggtcactga aaaatcttca attggactat gttgacctct atcttattca tttccagt 360
 tctgtaaagc caggtgagga agtgatcca aaagatgaaa atggaaaaat actatttgac 420
 acagtggatc tctgtgccac rtgggaggcc atggagaagt gtaaaagatgc aggattggcc 480
 aagtcacatc ggggtgtccaa cttcaaccac aggtctgtgg agatgatcct 530

<210> 88
 <211> 529
 <212> DNA
 <213> Homo sapien

<400> 88
 acctgagcta agaaggataa ttgtcttttg gtaactaggt ctacagggtt acatttttct 60
 gtgttacact caaggataaa ggcaaaatca attttgtaat ttgttttagaa gccagagttt 120

| | | | | | | |
|-------------|-------------|-------------|-------------|------------|-------------|-----|
| atctttttcta | taagttttaca | gccttttttct | tatatataca | gttattgcca | cctttgtgaa | 180 |
| catggcaagg | gacttttttta | caattttttat | tttatttttct | agtaccagcc | taggaattcg | 240 |
| gtagtacttc | atttgtattc | actgtcactt | tttctcatgt | tctaattata | aatgaccaa | 300 |
| atcaagattg | ctcaaaagg | taaatgatag | ccacagtatt | gctccctaaa | atatgcataa | 360 |
| agtagaaatt | cactgccttc | ccctcctgtc | catgaccttg | ggcacaggga | agttctgggtg | 420 |
| tcatagatat | cccgttttgt | gaggtagagc | tgtgcattaa | acttgcacat | gactggaacg | 480 |
| aagtatgagt | gcaactcaaa | tgtgttgaag | atactgcagt | catttttgt | | 529 |

<210> 89

<211> 547

<212> DNA

<213> Homo sapien

<400> 89

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| gtttatatat | atagcgaata | aatctagttg | tataaatttt | taaatgccgt | cagtagaaag | 60 |
| cacacaaggt | tatgattttt | ttaattactg | gcttctgatt | tctttcactt | ctgacccctt | 120 |
| tcctttttct | cagatgtagc | tgagtcttga | tcattttaag | acaacgatgg | gtagaatttt | 180 |
| gagattaatg | ttaattttcc | ctttttgtta | atttcagtc | cctctcacta | tgcttttgtc | 240 |
| cagaaggatc | aagaattcta | ccatcccttg | ggtctttgtg | tataaacaat | gttaaataaa | 300 |
| ggtagactca | gtctttaaga | tattagacag | tttttttagt | ccatgggatt | gtaaatataa | 360 |
| acattaactt | tcctataaga | atattttggc | tttgtaatct | atagcctcaa | attgggtattt | 420 |
| attatggatt | cactagacaa | acagctgttt | ccttattgtc | ttttttcttt | agtgtttctg | 480 |
| atttgctatc | agtagctgtt | tttaaagcca | tccaaggaaa | ataattattt | acagtttttg | 540 |
| aagtcac | | | | | | 547 |

<210> 90

<211> 528

<212> DNA

<213> Homo sapien

<400> 90

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| gagcagcaga | agctgtacag | caagatgatc | gtggggaacc | acaaggacag | gagccgctcc | 60 |
| tgagcctgcc | tccagctggc | tggggccacc | gtgcgggggtg | ccaacgggct | cagagctgga | 120 |
| gttgccgccc | ccgccccac | tgctgtgtcc | tttccagact | ccagggctcc | ccgggctgct | 180 |
| ctggatccca | ggactccggc | tttcgcccag | ccgcagcggg | atccctgtgc | acccggcgca | 240 |
| gcctaccctt | ggtggtctaa | acggatgctg | ctgggtgttg | cgaccagga | cgagatgect | 300 |
| tgtttctttt | acaataagtt | gttgaggaa | tgccattaaa | gtgaactccc | cacctttgca | 360 |
| cgctgtgcgg | gctgagtgg | tggggagatg | tggccatgg | cttgtgctag | agatggcggt | 420 |
| acaagagtct | gttatgcaag | cccgtgtgcc | agggatgtgc | tgggggcggc | caccgcctct | 480 |
| ccaggaaagg | cacagctgag | gcactgtggc | tggcttcggc | ctcaacat | | 528 |

<210> 91

<211> 547

<212> DNA

<213> Homo sapien

<400> 91

| | | | | | | |
|------------|-------------|-------------|-------------|-------------|------------|-----|
| atataccatt | taatacat | taatacat | taatacat | taatacat | taatacat | 60 |
| gacatataga | actttacaaa | catatgtcca | aggactctaa | attgagactc | ttccacatgt | 120 |
| acaatctcat | catcctgaag | cctataatga | agaaaaagat | ctagaaaactg | agttgtggag | 180 |
| ctgactctaa | tcaaatgtga | tgatttgaat | taraccmttt | ggscyttgra | ccttymtwrg | 240 |
| raaaawgrmc | cmaccttityt | taacmtgrac | cwccytmatc | tctagaagct | gggatggact | 300 |
| tactatyctk | gttwatattt | taaatackga | aagggtgctat | gcttctgtta | ttattccaag | 360 |
| actggagata | ggcagggcta | aaaagggtatt | attatttttc | ctttaatgat | ggtgctaaaa | 420 |

```
<210> 92
<211> 527
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(527)
<223> n = A,T,C or G
```

```
<210> 93
<211> 531
<212> DNA
<213> Homo sapien
```

```
<210> 94
<211> 547
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1) ... (547)
<223> n = A,T,C or G
```

<400> 94
gttaaacatg gtctgcgtgc cttaagagag acgcttcctg cagaacagga cctgactaca 60
aagaatgttt ccattggaat tgttggtaaa gacttggagt ttacaatcta tgatgatgat 120

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| gatgtgtctc | cattcctgga | aggtcttgaa | gaaagaccac | agagaaaggc | acagcctgct | 180 |
| caacctgctg | atgaacctgc | agaaaaggct | gatgaaccaa | tggaacatta | agtgataagc | 240 |
| cagtctatat | atgtattatc | aaatatgtaa | gaatacaggc | accacatact | gatgacaata | 300 |
| atctatactt | tgaacaaaaa | gttgcagagt | ggtggaatgc | tatgttttag | gaatcagtc | 360 |
| agatgtgagt | tttttccaag | caacctcact | gaaacctata | taatggaata | catttttctt | 420 |
| tgaagggtc | tgtataatca | ttttctagaa | agtatgggta | tctatactaa | tgtttttata | 480 |
| tgaagaacat | aggtgtcttt | gtggttttta | agacaactgt | gaaataaaat | tgtttcaccg | 540 |
| cctggtg | | | | | | 547 |

<210> 95

<211> 1265

<212> DNA

<213> Homo sapien

<400> 95

| | | | | | | |
|-------------|------------|-------------|-------------|------------|------------|------|
| gtgggtcaagc | agtgattttt | ctgggactgc | agaagttcct | gctgtgceca | acctttatta | 60 |
| ctaactggga | aagaccagg | gagactggga | tgggtcctatg | attctacata | cagaactcat | 120 |
| ccaagaaagg | aggaagct | gatttttgtg | aacgtcgcta | cttgtgcctg | aactaactct | 180 |
| caggcacatt | agtcagaaaa | tactacctat | ggttactccc | ccaggttcct | aaaagtaaag | 240 |
| cttttagaggc | caccaaattg | gcaattgaag | ctggtctccg | ccatattgat | tctgctcatt | 300 |
| tatacaataa | tgaggagcag | gttggactgg | ccatccgaag | caagattgca | gatggcagtg | 360 |
| tgaagagaga | agacatattc | tacacttcaa | agctttgggtg | caattcccat | cgaccagagt | 420 |
| tggtccgacc | agccttgga | aggtcactga | aaaatcttca | attggattat | gttgacctct | 480 |
| accttattca | ttttccagtg | tctgtaaagc | cagggtgagga | agtgatccca | aaagatgaaa | 540 |
| atggaataat | actatttgac | acagtggatc | tctgtgccac | gtgggaggcc | gtggagaagt | 600 |
| gtaaagatgc | aggattggcc | aagtccatcg | gggtgtccaa | cttcaaccgc | aggcagctgg | 660 |
| agatgatcct | caacaagcca | gggtcaagt | acaagcctgt | ctgcaaccag | gtggaatgtc | 720 |
| atccttactt | caaccagaga | aaactgctgg | atttctgcaa | gtcaaaagac | attgttctgg | 780 |
| ttgcctatag | tgctctggga | tcccaccgag | aagaaccatg | ggtggaccgg | aactccccgg | 840 |
| tgctcttgga | ggaccagtc | ctttgtgcct | tggaacaaaa | gcacaagcga | accccagccc | 900 |
| tgattgccct | gcgctaccag | ctrcagcgtg | gggttgtggt | cctggccaag | agctacaatg | 960 |
| agcagcgc | cagacagaac | gtgcaggttt | ttgagttcca | gttgactgca | gaggacatga | 1020 |
| aagccataga | tggcctaaac | agaaatgtgc | gatatttgac | ccttgatatt | tttgctggcc | 1080 |
| cccctaatta | tccattttct | gatgaatatt | aacatggagg | gcattgcatg | aggtctgcca | 1140 |
| gaaggccctg | cgtgtggatg | gtgacacaga | ggatggctct | atgctggtga | ctggacacat | 1200 |
| cgctctgggt | taaatctctc | ctgcttgggtg | atttcagcaa | gctacagcaa | agcccattgg | 1260 |
| ccaga | | | | | | 1265 |

<210> 96

<211> 568

<212> DNA

<213> Homo sapien

<400> 96

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| ccagtgtggt | ggaattcggt | ttaattacaa | aatttgatca | cgatcatatt | gtagtctctc | 60 |
| aaagtgtctc | agaaattgtc | agtggtttac | atgaagtggc | catgggtgtc | tggagcacc | 120 |
| tgaactgtta | tcaaagttgt | acatatttcc | aaacattttt | aaaatgaaaa | ggcactctcg | 180 |
| tggtctctctc | actctgtgca | ctttgctgtt | ggtgtgacaa | ggcattttaa | gatgtttctg | 240 |
| gcattttctt | tttatttgta | aggtgggtgg | aactatgggt | attggctaga | aatcctgagt | 300 |
| tttcaactgt | atatatctat | agtttgtaaa | aagaacaaaa | caaccgagac | aaacccttga | 360 |
| tgctccttgc | tggcggttga | ggctgtgggg | aagatgcctt | ttgggagagg | ctgtagctca | 420 |
| gggctgtcac | tgtgaggctg | gacctgttga | ctctgcaggg | ggcatccatt | tagcttcagg | 480 |
| ttgtcttggt | tctgtatata | gtgacatagc | attctgctgc | catcttagct | gtggacaaaag | 540 |
| gggggtcagc | tggcatgaga | atattttt | | | | 568 |

<400> 97

```
<210> 98
<211> 547
<212> DNA
<213> Homo sapien
```

<400> 98

| | | | | | | |
|-------------|------------|------------|------------|-------------|------------|-----|
| tactgggtgc | caagctatgt | gccaggcact | ttacatgtat | tgattttaaca | cttaacagcc | 60 |
| actctatatt | attccctttt | tacagatgag | gcaatttaag | ctcaaagcat | ttaagtagac | 120 |
| aaccaacctt | gaatcacata | gcaaatgaca | gaagccagag | gcctcccaag | tctctctaac | 180 |
| tccaaaccct | atgcttactc | tactatatca | cactaccttg | caataggaca | aagggaatat | 240 |
| gtggtaaaact | atgttccag | catctaaaag | ccaggagtgg | ttttcatttt | tctttaagaa | 300 |
| gatgatagt | tgatttgaaa | catatctgaa | tttcagaaga | ggggactttt | aaaaattgcc | 360 |
| actcataagg | aaagaaagaa | ctttttcaca | tatttttgaa | agaaacgatg | gtgagaagat | 420 |
| attcttgata | atagagatat | gctaacattt | gctttgggtg | ttttgtaggt | tagatttttt | 480 |
| tgggtgtgtac | tttataggct | tgcatattgc | ttacttttaa | cagctgaagt | tctaagtaag | 540 |
| agtgttc | | | | | | 547 |

```
<210> 99
<211> 122
<212> DNA
<213> Homo sapien
```

<400> 99

```

cagcctttct gtcacatct ccacagccca cccatcccct gagcacacta accacctcat      60
gcaggcccca cctgccaaata gtaataaagc aatgtcactt ttttaaaaca aaaaaaaaaa    120
aa                                                122

```

```
<210> 100
<211> 449
<212> DNA
<213> Homo sapien
```

<400> 100

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ctgacggcctt | tgctgtccca | gagccgccta | aacgcaagaa | aagtcgatgg | gacagttaga | 60 |
| ggggatgtgc | taaagcgtga | aatcagttgt | ccttaatttt | tagaaagatt | ttggtaacta | 120 |
| gggtgtctcaq | ggctgggttq | gggtccaaag | tgtaaggacc | ccctgccctt | agtggagagc | 180 |

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| tggagcttgg | agacattacc | ccttcatcag | aaggaatttt | cggatgtttt | cttgggaagc | 240 |
| tgttttggtc | cttgggaagca | gtgagagctg | ggaagcttct | tttggctcta | ggtgagttgt | 300 |
| catgcgggta | agttgaggtt | atcttgggat | aaagggctct | ctagggcaca | aaactcactc | 360 |
| taggtttata | ttgtatgtag | cttatatttt | ttactaaggt | gtcaccttat | aagcatctat | 420 |
| aaattgagtt | ctttttctta | gttgtatgg | | | | 449 |

<210> 101
 <211> 131
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| <400> 101 | | | | | | |
| ccatgttctc | tcttgactac | gcatatgtga | gatttgcccc | tccgccccgc | tcgtgatagc | 60 |
| catccagatc | ttttacctgg | ccctgtcttg | gagaatctgt | tttcaatctc | cactgattgc | 120 |
| ccccttgctg | g | | | | | 131 |

<210> 102
 <211> 199
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| <400> 102 | | | | | | |
| ctgctgcgcc | tgatgctggg | acagccccgc | tcccagatgt | aaagaacgcg | acttccacaa | 60 |
| acctggattt | tttatgtaca | accctgaccg | tgaccgtttg | ctatatccct | ttttctatga | 120 |
| aataatgtga | atgataataa | aacagctttg | acttgaaaaa | aaaaaaaaaa | aaaaaaaaaa | 180 |
| aaaaaaaaaa | aaaaaaaaaa | | | | | 199 |

<210> 103
 <211> 321
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|-------------|-------------|-----|
| <400> 103 | | | | | | |
| tttttttaggt | ttttaaactt | tttatttgca | tattaaaaaa | attgtgcatt | ccaataatta | 60 |
| aaatcattttg | aacaaaaaaa | aatggcactc | tgattaaact | gcattacagc | ctgcaggaca | 120 |
| ccttgggccca | gcttggtttt | actctagatt | tactgtcgt | cccacccccca | cttctttcac | 180 |
| cccactttttt | ccttcaccaa | catgcaaagt | ctttccttcc | ctgccaccca | gataatatag | 240 |
| acagatggga | aaggcaggcg | cggccttcgt | tgtcagtagt | tctttgatgt | gaaagggggca | 300 |
| gcacagtcac | ttaaacttga | t | | | | 321 |

<210> 104
 <211> 309
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|-------------|------------|-------------|-------------|-------------|------------|-----|
| <400> 104 | | | | | | |
| tttttttttt | tttttatttt | tttttttgca | tcaaaaaaact | ttattttccat | ttggcccaag | 60 |
| gcttggttagg | atagttaaaa | aagctgccta | ttggctggag | ggagaggctt | aggcaaaacc | 120 |
| cctattactt | tgcaaggggc | ccttcaaaaag | tctctgggct | tctatttcaa | cgcgatgat | 180 |
| gtggctctgg | aaggcgtgag | ccactttttc | cgggaactgg | ccaaggaaaa | gcccaggggc | 240 |
| tacaaccgtt | tcctgaaaat | gcaaaaccag | cggggcggcc | gcgctctttt | ccaggacatc | 300 |
| aaaaagcca | | | | | | 309 |

<210> 105

<211> 591
 <212> DNA
 <213> Homo sapien

<400> 105
 cttatttctg catgggtcgg agagtgggag ggactgcttt actgagttat agtgaatgta 60
 gttttaacct aagcgctca catgactaac tctcatcca tcaagaatga gctcagctct 120
 cacttcccca ctctcacc cctgtaaaag taacctttct ccaagggtat gcttcaacag 180
 gaatagctaa catatttaa attgtggcac gtaagtatct tggatatatt ggctcattga 240
 atcctcacac ctactatttt acagagatgc cagtggggct tgagattgaa tcacttgccc 300
 aggctccac tgctggtaaa cagtagaggg ggctcctgac ccatcagtct ggcttgacaa 360
 cccattccct caactgcgga tcccggaatt ccttatcacc ctgttgattt ctccataggc 420
 tgtggtaaca tttgttgcac gaatggaccg ttgaaatagg gcctggcagg gagaaattca 480
 ggaaatgaat gaatggttct tccctggcag cctttgatga cttacaagcc ccttcaaggg 540
 ggaaagccat ttttctccct gggactcctt gaaagcccg gaggcctgcc t 591

<210> 106
 <211> 450
 <212> DNA
 <213> Homo sapien

<400> 106
 ctgccactcc tgctctgct accccgaaac cggagaggga gctcaataat aacacaggtc 60
 ccactaaact aattaagggt ttggcataac ctgtcattga attcaagtgt ccaacaactg 120
 tttgcttaaa atatcattag acctaataatt tttttcaaag gcacaaagtt taaacatggg 180
 gggggcgggt gttgagaggg gtctgggata cccttaaacc caaaaaagtg atttggtccc 240
 ccttgcccag aagggtgact gttccactgg gcctgtcacc acaggacatt ttccatgaca 300
 agcactcacc ttcttgggga aggggcatca gggtggcaca ggaaaggccc aagtgagggg 360
 ccactctgta cattaatact ttggtgatta atgtttgggg agaggcagga ttctcaccca 420
 cctttttgac ttcaaacact ctcaactcaag 450

<210> 107
 <211> 116
 <212> DNA
 <213> Homo sapien

<400> 107
 tcgacgaaag ttactgtcac tcagttgtaa atccatcagc ttttcacctg ttaaaaattt 60
 tgcaaaatat acatgttctc ctctgtttt caattcttcc atcttttttc ttgagg 116

<210> 108
 <211> 291
 <212> DNA
 <213> Homo sapien

<400> 108
 ctgctogaag ttgtcaaaac ccacgtgcag ggcaatggag agtccgatgg ccgaccacag 60
 cgagtagcgt cctcccaccc aatcccagaa ctgaacatg ttttgagggt caattccaaa 120
 ctcccttact ttggttgtgt tagtagacag ggcaacaaag tgcttcgcca ctgcagtagg 180
 atccttggcc gcctggagaa accactcctt cgccgtctct gcattcgtga tggctctctg 240
 ggtagtaaag gtcttgaggg caatgatgaa cagggaggac tcgggggttca g 291

<210> 109
 <211> 662

<212> DNA
<213> Homo sapien

<400> 109
gctgtttcca cagtacgcct gcctcacacc ttgcatgacg ccaacatcac catcattgag 60
caccagaagt gtgagaacgc ctaccccggc aacatcacag acaccatggg gtgtgccagc 120
gtgcaggaag ggggcaagga ctctgccag ggtgactccg ggggcccctct ggtctgtaac 180
cagtctcttc aaggcattat ctctggggc caggatccgt gtgcgatcac ccgaaagcct 240
ggtgtctaca cgaaagtctg caaatatgtg gactggatcc aggagacgat gaagaacaat 300
tagactggac ccaccacca cagcccatca cctccatttt ccacttgggtg tttgggtcct 360
gttactctctg ttaataagaa accctaagcc aagaccctct acgaacattc tttgggcctc 420
ctggactaca ggagatgctg tcaacttaata atcaacctgg gggtcgaaat cagtgaagacc 480
tggattcaaa ttctgccttg aaatatgtg actctgggaa tgacaacacc tggtttgctc 540
tctgttgat cccagcccc aaaagacagc tcctggacct tgccccgggg cggcccgcctc 600
ggaaaggggg cgaaatttct tcaagaatat ttccatttcc acaaacttgg ggccgggggc 660
cc 662

<210> 110
<211> 323
<212> DNA
<213> Homo sapien

<400> 110
tcctgtgaaa cagcccattt tcctacctac tgtgggttgc tgctcaggag gaacgatata 60
cgccaatata agcaggaaat ctgcagctcc tctgctatgt gcctcagaac actttcaatt 120
tttctgggtca atgctctgat taggtatcat acataaaaagc cagcatatta gtttaaatct 180
ctaacaaaaa actatatttt ccaaagtcac tatcatttgg gccaatataag tgatcttttc 240
gtgctttgtt gagcttcac tttagggcac ctcttcttcc ttcccattca tgaagtccgg 300
catttccatg tgcaaattta cag 323

<210> 111
<211> 336
<212> DNA
<213> Homo sapien

<400> 111
tccagtgcgc tccagcctta tctaggaaag gaggagtggg tgtagccgtg cagcaagatt 60
ggggcctccc ccatcccagc ttctccacca tcccagcaag tcaggatata agacagtcct 120
cccctgaccc tcccccttgt agatatcaat tccataacag agccaaatac tctatatcta 180
tagtcacagc cctgtacagc atttttcata agttatatag taaatggctc gcatgatttg 240
tgcttctagt gctctcattt ggaaatgagg caggcttctt ctatgaaatg taaagaaaga 300
aaccactttg tatattttgt aataccacct ctgtgg 336

<210> 112
<211> 218
<212> DNA
<213> Homo sapien

<400> 112
tttttttttt tttttttttt tccagtcagg agtattttta atcactgtct acagagacac 60
ctacatacac acacgggtgg ggaatgaacc caaagttttt aggtgaagtc tctcagggcc 120
caccocgtgc cacagacctt cctcggttgc agagattctg ggcaaagcat ccgtgctctc 180
atgagattat cctggggaga tttagaagaa ttttgtgg 218

<210> 113
 <211> 533
 <212> DNA
 <213> Homo sapien

<400> 113
 ctgcaccgac agttgcatg aaagtctctaa tctcttccct cctcctgttg ctgccactaa 60
 tgctgatgtc catggtctct agcagcctga atccaggggt cgccagaggc cacagggacc 120
 gaggccaggc ttctaggaga tggctccaga aaggcggcca agaattgtgag tgcaaagatt 180
 ggttcctgag agccccgaga agaaaattca tgacagtgtc tgggctgcca aagaagcagt 240
 gcccctgtga tcatttcaag ggcaatgtga agaaaacaag acaccaaaagg caccacagaa 300
 agccaaacaa gcatcccaga gcctgccagc aattttctcaa acaatgtcag ctaagaagct 360
 ttgctctgcc tttgtaggag ctctgagcgc ccactcttcc aattaaacat tctcagccaa 420
 gaagacagtg agcacaccta ccagacactc ttcttctccc acctcactct cccactgtac 480
 ccaccctctaa atcattccag tgctctcaaa aagcatgttt ttcaagatct aaa 533

<210> 114
 <211> 261
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(261)
 <223> n = A,T,C or G

<400> 114
 ccatatctgc tcggcgctac ttctttcttg gattgatcct gantgatgca ttggcgatgc 60
 ctttgagaaa ggacatgtga tgtgatggtc ttcacgttcc acatgtactc gggcaaatag 120
 ggggacaaac tgaagttaaa caggctgaaa ctagaggagc tgctgaccct ggagctgacc 180
 actttcttgg ggaaaaggac acatgaaggt gctttgcaaa agctgatgag caatctggac 240
 accaacatag gacaacaacg t 261

<210> 115
 <211> 267
 <212> DNA
 <213> Homo sapien

<400> 115
 cctctcctgt ggggttccaga ccctgttcca gcaacaattg ctgggacacc tgggccgact 60
 gctccacctc gccaggccct ggccctctcc atctcagccc tgacagccac ccagtataaa 120
 acacagcagg cttcctaagc aatgtgacgc accagagggg tgggtggtaca cgttccctt 180
 gaagtcactc gaaaattaga gaacagattt gcctcatagc tgaagagaga ccctattcca 240
 agcatgaatg gccttgacaa tgttcct 267

<210> 116
 <211> 239
 <212> DNA
 <213> Homo sapien

<400> 116
 ctgatgacct ggggtctagt gaaaatgcag ggtcagattc agtgggtctg ggggtctgaat 60
 ctctaaggcg ctgccaaagt atgctgatgc tcctggcttg tggaccacc cgtgtatagc 120
 aaagctctag actaggaggt ctcaaccttg gctgcacaga attatctggg gagtttttaa 180

```
<210> 117
<211> 168
<212> DNA
<213> Homo sapien
```

```
<210> 118
<211> 150
<212> DNA
<213> Homo sapien
```

```
<210> 119
<211> 154
<212> DNA
<213> Homo sapien
```

```
<210> 120
<211> 314
<212> DNA
<213> Homo sapien
```

```
<210> 121
<211> 601
<212> DNA
<213> Homo sapien
```

```

<400> 121
aaaaaaaaacc taattcattg aagtaataac caaataatTT tcaatcttga ttcaactgtg      60
attcaaatct tacaccattt gccctttcta tgaatttatg tataaaatTT tttaagagtc      120
agagttttttt tttcttgatt aattggatgt atttcacaga atttccaact gctcacgtta      180

```

```

gttttcttcc ttttagagtt gatctctcta atgtattaga tcttcatgcc tttgatagtc      240
tctctggaat aagtttgcag aaaaaacttc agcatgtgcc aggaacacaa cctcaccttg      300
atcagagtat tgtacaatca catttgacgt accaggaaat gcaaaggaag aacatcttaa      360
tatgtttatt cagaatcttc tgtgggaaaa gaatgtgaga aacaaggaca atcactgcat      420
ggaggtcata aggctgaagg gattgggtgtc aatcaacgac aaatcacaac aagtgattgt      480
ccagggtgtc catgagctct gtgatctgga ggagactcca gtgagctgga aggatgacac      540
tgagagaaca aatcgattgg tcctcattgg cagaaattta gataaggata tccttaaaca      600
g                                                                                   601

```

<210> 122

<211> 486

<212> DNA

<213> Homo sapien

<400> 122

```

ctgttttctaa ttgcttttgt gactgttacc ttttagttca tgccccccca aagagctaaa      60
tttcacattt ttacctacaa aattgatttt taattcctgc aaataattta ccattatgag      120
ctacaagggtg ggcaacagcg cctgaggatc taattttatg catattactc ccaagtattt      180
taacacttgt tggagaagca atatctggat caataaaaca ctgtcccatc aaccatttga      240
gtggggagag ggagaagctc ttctgtaagt aagattcttg caagctcttt gaaatgagtc      300
ttctttccca cagattttct ctactctttc aatacaaaaca gataggagaa gaggggaatag      360
aaacctggag gaacttgaat atttttgttc tagatagaga tacagttatt gaaaaggaaa      420
cctagaaagt agtcacacgt cgcttattta ggccagaagt aattgtactg ggcaaaaatt      480
tcactt                                                                                   486

```

<210> 123

<211> 239

<212> DNA

<213> Homo sapien

<400> 123

```

ctgggtgggtc ttttttttct ctcagagctc aagcctgtag tgctgatgt catttctttc      60
aagttgcccc cagtatctcc acttaaaacta ggctagtaac caaaataatg tggaccttct      120
ttaggaaaca gtgtgggaga ataggagtcc agccgtaaga taaactggaa atatttgggc      180
gtcttgtagc tggctacgca ccacctcagt gttgttccta cataaacaag gcccccttt      239

```

<210> 124

<211> 610

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (610)

<223> n = A,T,C or G

<400> 124

```

ccanccaagt cnttgatgat cactgaccen cgcgcgcctg ctggaccaag gtggtgcgg      60
ggaaatcgcc acngngcttt cggttttctt ggtgaaggaa tacaccgcgc cgacagcagg      120
ttttcagtca gggtcaggga ctggtgcttg cgcgcgaaaa tcaccgttac gccgaggttc      180
aggccggtca tgatcgccgg tgcaatgcc gaggcttcga tggtgacgat cttggtgatg      240
cccgaatcct tgaacaacgc agcgaattca tcaccgatca gtttcatcag cgccggggtcg      300
atctggtggt tcagaaaggc gtcgaccttg agtacctgat cggaaagcac gatgccttct      360
tcgcgaatth tcttgtgcag tgcttcacg aaagcttcct ctggtggcgc aacacgcgcc      420

```

gaaagtagat taaaaagtag tcgattctag cgctttaaca tcgcgcgtat atccgccagg 480
 gcggtattgc cgcgaacggc tttgacttcg gttggtgtgt cgtcgttgcc ttcccatgcc 540
 aggtcatccg gcggcagttc gtcaaggaac cggctggggg cacaatcaat gatctcgccg 600
 tactgcttgc 610

<210> 125
 <211> 196
 <212> DNA
 <213> Homo sapien

<400> 125
 ctatagggct cgagcggccg cccgggcagg taaaaaatca gcccctaatt tctccatgtt 60
 tacacttcaa tctgcaggct tcttaaagtg acagtatcct taacctgccca ccagtgtcca 120
 cctccggcc cccgtcttgt aaaaagggga ggagaattag ccaaacactg taagctttta 180
 agaagaacaa agtttt 196

<210> 126
 <211> 247
 <212> DNA
 <213> Homo sapien

<400> 126
 aaattagtta aaaaaatgca ttcctcattt gatatagcca cattccaaat gcttaaaagc 60
 cgcattgtatc tagtgactac catactggag agtacaaata tagaacttta cccgtcactg 120
 cagacagttc tggttgattg tgcagcattg gacaatatat acagtttgcc tgtatatgag 180
 aaagagagag agagagagag tgtgtgtgtg tgtgtgtgtg tgaagtgcaa taaggctgac 240
 aggcac 247

<210> 127
 <211> 590
 <212> DNA
 <213> Homo sapien

<400> 127
 cctccacggc atggcgcaat tgttgttcag gggccgccag gttgctgcc atgccgatgt 60
 agatacgttc cactgtgctta ctgccagac gcactcgaag cgtcgccagc gctacgtttg 120
 cgcttgctgc cactgctgcg gcgacgctt ttcgggccat cgccggtggc ttgcctttg 180
 ctgctgagct ctttgatcat ctgcggcgc tggctgtcgt tggcgtcctg gtagtcggtc 240
 caccactcgc caaggccgtc ggtctgttcg ccggcgcttt cacgcagcag caggaagtca 300
 tagcccgga cgggaagcgcg ggttgctccag caacaggtcg gcacgtttgc cgctgcggcg 360
 tggcaggcgc tcctgcatgt cccagatttc acggatcggc atggtgaagc gtttcgggat 420
 ggcgatgcgc tggcattgct cggcgatcag ctcgtagca gcttcctgca tggctggaat 480
 tgccggcatg ccacggctct gcaggcgcg gacgcgttcc gaaagcgcgg gccacaacag 540
 ggccggcaaag aggaacgccg gggtgaccgg tttgttctgc ttgatgcgca 590

<210> 128
 <211> 361
 <212> DNA
 <213> Homo sapien

<400> 128
 ctgcccattg aaaccctcca ggagctgctg gacctgcaca ggaccagtga gagggaggcc 60
 attgaagtct tcatgaaaaa ctctttcaag gatgtaacca aagtttccag aaagaattgg 120
 agactctact agatgcaaaa cagaatgaca tttgtaaacy gaacctggaa gcatcctcgg 180

<222> (1)...(305)

<223> n = A,T,C or G

<400> 131

```

aaacacatac gaatanttna actgtgatta tgaagtgaca gccggctaaa tatgtcttgt      60
atthttctctc ttcttttttt tgctaactca tcctttattc cattcctgct tccatggtaa      120
tgcaggctca aataaattac taggatacaa gattacttca agcctctttt ctgtggaact      180
cataatatga taagcatttg ttacaagatt gcctgtagtt gtttagggga caaattatat      240
tagggaaaaga aagtctttct ttagttgggt aaattttcta ttataattgg gtactaaatt      300
tattt                                           305

```

<210> 132

<211> 545

<212> DNA

<213> Homo sapien

<400> 132

```

aaacaatgct acactcattt ttggcaaagt gctgtattgt tcagtctgtg tacaaaactg      60
accatctatg aaccaatcag tataaaaaat ttctataaaa acaaaattta gacagcggct      120
caagaaaaca agctgccatt tatgcataga ttgatgtaca gtaacctaac caaatgtccc      180
ttttgaattt tcaagttact gaaaaaaaaat gtgtcgagaa acacattaag aaggcacatg      240
tacagtctac aatactcttc agtctcccta actcatgccc tgcccctata aaggaaatat      300
gttcacaatt ttacttgaga aaaaaaaaca aagccactta aaaaaaaaaa aacacacacg      360
caattattaa agttcaaaat ctctggagga aaatacaagc aaaaccactc atacactcca      420
agcctgaaac acacatctaa cctccccagg tactggtttg gttttcagag gtccacctag      480
aaaacaaatc taaaacttca ggcaaaacag agcaaaactg gacatttaac aattacacaa      540
ttttt                                           545

```

<210> 133

<211> 330

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(330)

<223> n = A,T,C or G

<400> 133

```

aatattttatt actaatatct tataatgttt tgtggnacca tggcatacct tgggtactat      60
tgtaacanat agttcaggaa accctactat aaggtttatc aaatgggtctc ataaacagtt      120
acttattcaa gcacgccaaa gctcagtga aagtattttt cacccttact ctttctcgtg      180
tcattcaaag agaagttttg atgtagtgta tttatttgta gggagtaatg aacagatcca      240
tttcacagta gactttgtgc tctaggtgat gcagctaatt gccccagttt ggaaaacatg      300
gacttggtatg aattgtcttt tgtttgggac                                           330

```

<210> 134

<211> 627

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(627)

<223> n = A,T,C or G

<400> 134

```

aaatattact tcaaatacat tttaaagctc aacaaacttg tgttgaactg aattgcagat      60
cctgaactct atttgaaaat acatcatgaa acagaaaanc ccattccaaa tgaaaatgat      120
agtgccttgt tgggggtggg aatgaggcgg ggagactaaa tcactattaa cagacttctt      180
ttcccaatgc aatttgtcaa aagttcaaaa gttctgaaat gtactaaatc ttaagcaaatt      240
taaattcatg atattactaa aactttttta atagtgcaat gacttatcaa gttatagtgg      300
ctgcattaag aacaaattat tgtgtgaaat acctgtataa acacaaaata caattaaata      360
tttctttaca aaaagctgag cattacgcac aatagtggaa tgtctttcat taggtgtatt      420
ttttaaagat taacaaaagt aacatttcct aaaatgtata catgtgccat atttttgcaa      480
acatgcctga gaatgtattt aaaacatttc tgtagtaaga gtttgcaaga acttcacaaa      540
cctgcaataa aaatgcatct ttttaaaaag gtgaaaatgg catctccaca ctgcaacaat      600
tcaaaaagtg cagcatccct aatctttt                                     627

```

<210> 135

<211> 277

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(277)

<223> n = A,T,C or G

<400> 135

```

aaaatcaaatt atattatttg ttaaaaatca gcttgtttca ttacnggaaa ttacaccagt      60
ccgttctatt tactttcaaa ccatattcaa ctctcaact ttcaaacatg taatcaacta      120
atttcaaaaag ggaaaaggta ccctttataa aggagagatc tgtaaagaca ccaagaaatc      180
aaaattaata tcacttaata attaagtgga taacacatgc ctccaatac agtgcagtga      240
gaaacacaaa acatcaattc ccgcgtactc tgcgttg                                     277

```

<210> 136

<211> 486

<212> DNA

<213> Homo sapien

<400> 136

```

aaaacagaat gaattcattg ttacagttac agaagtcaga agcccaaata cagtctgcct      60
gaaccaaagc cagggtcagc aagggttcctt tccactgttt tgccaacttc tagaggccac      120
ctgtattcct tgggttcattg cccctctctt catcatcaaa taatcagcat agctttatga      180
cattggcagc tctgattttg ctcttttgcc ttcctcttat gtagaccctt gtaattacat      240
tgggtacacc cagataaccc caaataatct ccctatctca agattcttaa tgtaattata      300
ttgggaaagt cccttttgct atataagata acatagcaat ggattccaag gattagtatg      360
tgagtttctt ttgaggggct ataattaacc ctaccacaat atggaaatgt ctattgtttt      420
tctatgtacc agaaataaga cattaggatg tgaaattaat aacataacac cacttacggc      480
atcacc                                     486

```

<210> 137

<211> 552

<212> DNA

<213> Homo sapien

<220>


```

acattggtgg cacttgaact gagtgcaaac cacaacattc ttcagattgt ggatgtgtgt      60
catgacgtag aaaaggatga aaaacttatt cgtctaattg aagagatcat gagtgagaag      120
gagaataaaa ccattgtttt tgtggaaacc aaaagaagat gtgatgagct taccagaaaa      180
atgaggagag atgggtggcc tgccatgggt atccatgggt acaagagtca acaagagcgt      240
gactgggttc taaatgaatt caaacatgga aaagctccta ttctgattgc tacagatgtg      300
gcctccagag ggctagggtta gtacaaactc gcattcatgg cttggtttcc cagaagatct      360
ccatttaact tttttaaaga aagtttattg ctttctttaa cctgcatttt ttctaagttt      420
tttttcgcat aaaggtgctg tctttgtggc aaggcctagg catgacaatc ggaggactcg      480
agggggatgg aggactagtg atccggctgg ctgcttccag tcgattagag aggtgaaaaa      540
gctgaacgtg tgcccantna atcttcaaaa aggcagaaac atatcacctt ntgccccnt      600
aaacttgttc tttttccgaa ggggaaaaaa aaaatggaaa      640

```

```

<210> 141
<211> 127
<212> DNA
<213> Homo sapien

```

```

<400> 141
aaaaatcaca cactgacaac acagaaatac gaaatgctag gaaaagtcta gcatatgaag      60
gaaaaacatg tcttatgcac tctaataataa ttttttcaat tagtataaag gcaaatgcgg      120
ttttttt      127

```

```

<210> 142
<211> 126
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (126)
<223> n = A,T,C or G

```

```

<400> 142
aaatatactc tggatgcntt caagtaatac taatcatttc atgngnaaaa gtcttttaat      60
aaacaaattc agagtaaaat taattgaaat atttataata catttgttac acagttattt      120
ccaata      126

```

```

<210> 143
<211> 730
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (730)
<223> n = A,T,C or G

```

```

<400> 143
gcaagttctg gagtgttcac ttctgagcct gaattccctc ccctgcaaaa tgggggaata      60
ccctcctcag agggtccttg cgagggtgag gggagatcag catggcaggt gtgctgggca      120
cggcagggcc tgggaagggc agatcctttc cccatccctg ccacaaacaa cccaaacctt      180
taaaggagag caatggcctt gtgtcaaaaa caaaaacaaa acaaaaccct gtcctaggag      240
actggggccc taattttctaa tagcaagcct ttatgagtcc ctaacactct actgggctga      300
gtatctcaca cgccagagga taacctgect tctgctcacc accaccccggt agtagttgtc      360

```

```

atttgtgtcca tttcacagat gaggcaaagg ctcagaagag tcatgtgtta aaccagcttc 420
tagagcccat gcaggagctg caggtgggga gaatcacctc taggtgctct tcccatggaa 480
tcctcaccct ccttgagtgg tcactcactc anctttccaa tgggtgtgtg acctttgacc 540
agctttcttt ccttntctgg gcctcagttt cccaccttgg acaaagtaag aggtctcttg 600
ggnttcangg tagttcttcc taacttcttt tccttttcat ttgagcatcc ttcttcattt 660
tttgccacct ctcttgatcat tacangcttt taccttcggc cgcgaaccac gcttaagggc 720
naaatttcca 730

```

```

<210> 144
<211> 485
<212> DNA
<213> Homo sapien

```

```

<400> 144
ctggtcagaa atgattctct tgtgacacca tcgccacaac aggcctcgggt ctgtcctccc 60
catatgttac ctgaagatgg agctaccttt cctctgtgtg gcattttgtc gcttatccag 120
tcttctactc gtagggcata ccagcagatc ttggatgtgc tggatgaaaa tcacctgtgt 180
tgcgtgggtg gtctgctgcc gccacttcta atcctcatca tgacaacgtc aggtatggca 240
tttcaaatat agatacaacc attgaaggaa cgtcagatga cctgactgtt gtagatgcag 300
cttcactaag acgacagata atcaaaactaa atagacgtct gcaacttctg gaagaggaga 360
acaaagaacg tgctaaaaga gaaatggtca tgtattcaat tactgtagct ttctggctgc 420
ttaatagctg gctctggttt cgccgctaga ggtaacatca gccctcaaaa atattgtctc 480
aacag 485

```

```

<210> 145
<211> 465
<212> DNA
<213> Homo sapien

```

```

<400> 145
ccaagacagc tcgtttctgg agagtatgag ggtgtgtttt cttattgtga aaggaaactac 60
cttctcttag agggtaggaa gaatgtgggtg tgtgtgtgtc tcataaagca accggacatt 120
ataggtgccc aggtcatcta taaaaacgat ccttgggctg tgtaaaaatg aagtggcttt 180
tcagtatcct ctttcacact tgctgcttcg ggagactatg caatgatggg aaggtgattg 240
cccctttatt tcattcagtg ccattggtccc tgttgttgta gtaatttatt tgtttagttc 300
atTTTTTTTT tcttaacagt caaggggaag agtgattcct cactactgctt tcaagctgga 360
ctgagccagt ctcatctctg gaaagaaatg ctgtgtccag aactcagcag ctccatctat 420
tttttccagt cgaaagaaac tgatcttttag gcagttttta cttgg 465

```

```

<210> 146
<211> 351
<212> DNA
<213> Homo sapien

```

```

<400> 146
ccagccgggg taatctgtat gtggcggact tgagctacga cgtgggcggc aagtgcctgt 60
ttgaccagat cagcggcgtg aagcttatgc caactcatcg ttgataaat ccgaggatca 120
gttcaagacg tcgcagcggg tgattttggg aacgtcgttt tcggtcagta aattgtgggt 180
agcgacggag tggttgatcg gcaagaatga tccgtatatt ggccgggagca gctataccga 240
gagcctgggg gctgggggga gtaaccagtg ggagaatcag ttatatatga acattgggta 300
ctacttctga cttaatatct ccagcgtttt aactggcctt atcgaggca a 351

```

```

<210> 147
<211> 654

```

<212> DNA
<213> Homo sapien

<400> 147
acttattttt aattactgaa tattttcttag acgttttggg acagatttta tgtaatcttt 60
ataagtatga tttctgaaga aaagcaaagt cattagtagt tttgccttaa acttgtagac 120
taaaccaagt attgtaaaat aaacagcgat aacagtgata gtttttaact ctatggcat 180
tgtatcactc tggaaaatgt ggagtagctg taataaatct actcctgtat tatgctttac 240
agtgcaggtc ttagtttttc ttttttctca tttcttttga aatggcatct cgaacaaagt 300
ccaccaatcc ctttacaaaa gaatgaactg ctctctctgtg tgtacttcat agaagggtga 360
atcgacaga ggcaggtag tgacagttat tcctgaaata caggagcaga gtacagtctg 420
ttgtggtttc ccgattccg cgcctagctc agccaattaa gcatgagaca taggccattg 480
agccacttag tagttatgcg agtggataga ttggtatgta agagggaaag aggtctgctg 540
taaagaacaa cacttgtttg tctgtgggga aagaaaagca gaatcttgag atgaaagttg 600
gcatacaaat aggatactat cgccagtagg ttatattaca aaacatttat cggg 654

<210> 148
<211> 539
<212> DNA
<213> Homo sapien

<400> 148
tgaatatcat gagggtgatt ttcacctgat tgcaaaactg ccatagtttg aaacactttt 60
tcaatttacc agacacactc tgtcaagact tcatatactt ccaacttgca agcctgtgtt 120
ttgccttctc caacctaaaa aggaaaagct ttaaacgatg aacttacatt ctattaaacc 180
atcagacttg agcttatcca tctgtttagc gtgaatgtac aaaccaggta catttccacc 240
aaacacatag aaaaatcttg tgcatacacag ttcagctaag ggtagtagga caatccttac 300
aatcctcctt ggatttcttt tttaagatgt caaagaagca ggtaagcaac attgttcatt 360
tgttactggg tgttctagat caaaccttca caagctatat atatagcttc atatgctata 420
gottacaaat ggggtaacaa agtaaaagaa aagaacaaat tatactttga cactttatag 480
tcaaagtata attaaaaaag aaatcctaca gtgggtaatg gagaaataga taatttttc 539

<210> 149
<211> 273
<212> DNA
<213> Homo sapien

<400> 149
tttttggtea ttctcctcaa ggagccgctg gatagtagtc ttgattgact tccaccttgc 60
ccctcatata gtccggtact aaggccaccg acatcccag gaacctccgg aaccacgacc 120
gccaagcaac tcgacccacg ataggtgggg cctacgctct cgaagttgat tggatgctcc 180
cgctacagg ggggggtaca gaaggacgt catttgtgac tggacgcgca agagctatac 240
tcagcagctt tcctctgtcc cagcccctag aac 273

<210> 150
<211> 200
<212> DNA
<213> Homo sapien

<400> 150
gtttttacta ccgtatggcc catttaaaag ggatgtgtac gccttacact ataaccctta 60
aaccacctag aaatatgaaa ctcaaactgc cactgacctc cctcaccaag ctccataaaa 120
gtaaaaaatt ataacaaccc ttattaacca aactgaacga acatatgggc gattgattca 180
ttgccccac aatcctaggg 200

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| tttttgtaa | cagtcttaat | aaataataaa | atggaataaa | gaaacccaaa | aaaaaagaaa | 120 |
| aagtttgat | gaaaattcat | ccctatttct | ttattttgga | ctaagtagtc | aaatttctac | 180 |
| tatattaata | ttatgtaagc | gacacccatt | taaattcact | ctctttgata | gaaaggtag | 240 |
| ttgattatca | cacctgctat | tttttcactg | ccaaaragac | tgcaataacc | tccctccatc | 300 |
| accctcaaaa | aacaaacaga | aaccatctga | ggcatagcca | ttgtttacat | attgtgtttg | 360 |
| tgtgcaccta | tctacaacgt | tctttcttct | aaggagttta | tctgccaata | ttttcggtt | 420 |
| cagcagcagc | gctcttcttg | acagactaag | agaaggatct | acagaaaagt | catctgatta | 480 |
| aggttttggg | tcaaattaaa | actctctgga | cagaatctct | tttcttctac | ttggatttct | 540 |
| gcaaacagaa | agcagattat | tctctgggca | caatagcgac | tctagaaacg | cttatgtttt | 600 |
| tcagactttg | gcagaacttg | ttaagaacag | catcatcata | atacatattg | acaaactcga | 660 |
| atttcagtgg | ctcttttctc | ccacatgatg | catgatgaaa | tttataaagg | tctgtttttac | 720 |
| ccccacagg | tcatttcttt | tgtgttccta | cagagccaat | aggcttcatt | taagtccaag | 780 |
| ttattatatt | aaccatccct | ttcactagac | tagagaactt | ctttttcatg | gtccatatcg | 840 |
| tga | | | | | | 843 |

<210> 155

<211> 674

<212> DNA

<213> Homo sapien

<400> 155

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tttctgtgca | gccccaggtt | tgctccagct | attcacaagc | agaatataac | acaagaaaaa | 60 |
| caattcatat | cccttaggga | aaaaagagga | tcaattcatc | actcaatatt | taatacagcc | 120 |
| aaaatgagct | gccaaaacaa | gcacacacac | aaatactgtg | aacagaaaaa | tacaagaaaa | 180 |
| tgactaagct | gggagctctg | acgggggatg | gacattgctt | aaagcactta | tcagtcccca | 240 |
| gaaaaaccaa | accaaaaaca | ttttttacga | tgccatggcc | tcattggccc | ctttaaaact | 300 |
| gttgatggta | acaaagggca | gggggtgggg | agagaaaaca | caatcactgc | tccctttttg | 360 |
| ctgcagctg | tgactgcacc | cctcacggca | cggcatgta | cacaactacc | acacaaggag | 420 |
| gaccaagtcc | ctctgctggt | ggcctcctaa | aaggcaaggc | ttgagttttg | gctgatgagc | 480 |
| aagttctctc | cgttaccaat | ccctgccaac | cagcactacc | atggctgaat | tgatctaccg | 540 |
| ttttctgag | taaactgtaa | ctggctacag | tttcggtaac | atggaaaaga | actcagctac | 600 |
| tacagccaac | tgcaatactt | caggaacccc | ctccatccct | ggggctcctc | actcctagt | 660 |
| catcttgatt | ggat | | | | | 674 |

<210> 156

<211> 671

<212> DNA

<213> Homo sapien

<400> 156

| | | | | | | |
|-------------|-------------|------------|------------|------------|-------------|-----|
| ccttttagtga | acacctttat | ctccatgtcc | ctcttagagc | ccagagagct | gcccataaggc | 60 |
| attttccaga | attctctcatg | tcacctagtt | caatttccat | taactcagat | cagccattgt | 120 |
| gattcaccat | ttgtcaggct | ctcaggttta | acaaaaccta | ctatcaccat | catccttcaa | 180 |
| cagccacagt | ctgaattgag | ccaacatttt | tttttctttg | agaaagaagt | gggctggggc | 240 |
| acaactttta | gtctgagggg | agctagtagt | cggcttgaca | attaaagcca | tccataacaa | 300 |
| cttttcttca | aatgtgttga | ctcctcaggg | gctaaactgc | tcttagctta | gaattatgct | 360 |
| ttactagaga | tctaccatat | aagtgggtta | atcactacca | tctgttaact | agttatatag | 420 |
| cttcagaca | tgaggagagc | atcaaacagg | gatggaagca | acccaagga | tatgcaagaa | 480 |
| gggcatgatg | aaccccttct | cctctggcag | gagaacaagg | ccaaccaagg | gacagactgg | 540 |
| aaagcactta | gatgtttaag | gaggagaaa | gggaagcttt | gaccagtcct | tgctttttgc | 600 |
| caagttcagc | cagttctccg | ctgcttgcaa | cctctagcgc | agtaacattt | tgagaattg | 660 |
| cagattttcc | c | | | | | 671 |

<210> 157

<211> 474
 <212> DNA
 <213> Homo sapien

<400> 157
 cgcgttcttt aattctttta gcttagaaag tcctttacac tacttaccta aaggccccaa 60
 agtaaaacac acactagtag taaggctagt gcatttcocct tctagcactc aaagaaagct 120
 taacattttt gacagtttgc aaataccgcc ttgtatttct gattcagcct tattcaaagt 180
 atcataataa aatattttatt aaatstatgt tgatctgcgt gcatttatga tctccagatt 240
 aacgttaggc ttctctgttg gcccctaact tggaggtgct tttttggatc cctcctcccg 300
 tgattcattg taatttcatt tcccttgta tggtctgac cagagaagat tctaaatatc 360
 tgccccaaa gccaaaatta tatcttttga aaagtgaat gaagagttga gtcastaatt 420
 tatttttagat attactgcct aaaacaattc cccaaaattt atggaagttg gagg 474

<210> 158
 <211> 584
 <212> DNA
 <213> Homo sapien

<400> 158
 ttggattctg cagttccaca tcattcactc cggcaaagga gagaacttgt aacaaagatg 60
 agtgccaagt ttagtcaatt taccctacct ggaatactat atacaactct ggggtctcatg 120
 tgtgttaaaa tacatacagt gaagctgagg aagagccact gaagtaaaaa gtattgttta 180
 caagttggaa aggatgtaaa aataatctaa agtatactaa gtcaggaata aaaggcagag 240
 ttaataaaaa tgtggctggg actgatagac gaaacagata tattttctaa atcctggaat 300
 aattattaaa aaattttaca tgtatcaatg gattccagac tccatatttt aagtttcaca 360
 actactgtca tttaaaacta taccttattg aacgtctccc actctcaata aattacccca 420
 aatcactctt ctccaaaacg taaatttgga acacactgac ttacaaaatt tgggcttaat 480
 ttataggatg ttgtggccct caaaaatatc attgtgggct aaacaaaata aattcttgaa 540
 acaattctaa aaatcaatca ttgtccaaaa tgaacttttt ctaa 584

<210> 159
 <211> 671
 <212> DNA
 <213> Homo sapien

<400> 159
 cctaatttta ttacttttct tgccactgct attattgata gaaatacaat taaataatta 60
 agatgaacca atccattgga agattactaa aattgtatct tccaatgcc tcctacagta 120
 agatttcttt ataattataa cccttggaaga caatttgaac tttattttaa tgttctgctc 180
 aaatctaaat ttccctctcc taggctgaag cctgatctaa ataaggaagt agttgggata 240
 tatccacagg ctgtcgaaca tggagctgca tctgagagac aggtggcagc aacccaaaagc 300
 aaagcaggga ctgagaacag gcaggttcca agagcaaaat ggaacttgaa agccaagtat 360
 ggttcactgt aaaggagaaa atatagaaat acggaactag aacacctggg ctgggatgtg 420
 gtaagcacco aaaatatagg aaaactgtat gaattcttgt gaagcagtaa actatgatag 480
 taatcatgtg acacatatga taacaaactc aaaacaggga aaagaggggc tttattcaat 540
 gctggagata agtgaaaaaa aaagtgaagt gtctcaagga cagaagttat catctcaaaa 600
 aggcataatca gctagatctc gcggaaacca tatgattatc ataattctag actctgttctg 660
 gtattacaaa g 671

<210> 160
 <211> 315
 <212> DNA
 <213> Homo sapien

ggtagacatc acctggatcc cccactctat tgcttacctt tttgttttgt aatttgatca 420
 gttcaagtta aaacaattta accaaaaact atgaatgttt atgatataat gaaatgattg 480
 ttaactttct tattgctttt tcacacacct ataaaagtaa ttttattact cccaagagaa 540
 atcactaaaag gcagaattac tagaggtaaa aataactagg gttggtacag tattactcag 600
 gagaagtcaa ggggagaaaa cttgtcccaa tgattcaaaa taattttggc atgggggggg 660
 ggaggggaaaa aaatttggct tccttt 686

<210> 164
 <211> 706
 <212> DNA
 <213> Homo sapien

<400> 164
 ttttttttgt ttcatttgct gcttaaaata aaaattataa attagattta aatggagcac 60
 taattataaa acagattgca agtaccacca tttgaaaaaa aaaaaaaaaa tcagtggatt 120
 tccataacac agaaaatgca tggacatgca tctacagtag agttaaaaaat ttcctgtgac 180
 taaaaaatta aaaactggaa tcaccagtag caaatgtata gtcaatggct atgacaagaa 240
 cagatcctgc cgagctcata aatgcaatta ttggcttttt tgctttataa aaaagacatt 300
 acatatttta ttgcattatt ctcttaataa aaaacatact accacgtagc tctccccatc 360
 cccattcttt gcttcagat ttttatagaa aataactgtt ttagtctggc cttggaaagt 420
 gaaccacca gcaccacct cacctactca ctcttcaatt caatatgcac atagcaaaag 480
 ccaacacttc aaatctcttg cccacatcaa aaaaagtagt ttcaggagaa aaacattaat 540
 accagttgaa taaaaataag ggcataaaag ctatgagaga gatagctctg ccatctgtct 600
 ctgggctaaa aatcaaggct aactattgcc tttggcacca caagggtcaa ggtccatggt 660
 tttattagaa aagtccccac aaaaaaatta aacccccctc acccca 706

<210> 165
 <211> 427
 <212> DNA
 <213> Homo sapien

<400> 165
 tyywgggcaa ttaggcagga gaaggaaata aagggtattc aattaggaaa agaggaagtc 60
 aaattgtccc tgtttgcaga cgacatgatt gtatatctag aaaaccccat tgtctcagcc 120
 caaaatctcc ttaagctgat aagcaacttc agcaamgtct caggatacaa aatcaatgta 180
 caaaaatcac aagcattctt atacaccaat aacagacaaa cagagagcca aatcatgag 240
 tgaactccca ttcacaactg cttcaaagag aataaaaatac ctaggaatcc aacttacaag 300
 ggatgtgaag gacctcttca aggagaacta caaaccactg ctcaaggaaa taaaagagga 360
 tacaacaaaa tggaagaaca ttccatgctc atgggtagga agaatacaata tggtgaaaat 420
 ggaaaaa 427

<210> 166
 <211> 124
 <212> DNA
 <213> Homo sapien

<400> 166
 accatgtttt ogttgtgtgt gagcagggaa gggaactttc ctgccttatt taaacctggg 60
 ccgaggattc gtggaatctg cttgatcaga gactctgagg ccaaaaacgc atcatacttc 120
 ttgg 124

<210> 167
 <211> 232
 <212> DNA

<213> Homo sapien

<400> 167

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tctgcatagc | aaatatgatt | taagaattta | acatcattat | ttgatcacao | gcgtaaatat | 60 |
| gtcaccataa | ataaatgtaa | attcattgta | caaaaattcc | caacaactct | taatacaaat | 120 |
| atggtacatt | tgacagtttc | tgaaacagat | tattttttaa | acttttttaa | acctaagctt | 180 |
| tatttttttc | ctggttatta | gacacacaca | aaaaaataa | aaagaggctg | gg | 232 |

<210> 168

<211> 677

<212> DNA

<213> Homo sapien

<400> 168

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| tttcacaatt | aaccaacatg | caaaaattct | cagactaaac | actgagaaat | tcttcataca | 60 |
| atgcatttgc | caccttattg | cattttttaa | atctttattc | tatagtgaat | tggtattccc | 120 |
| aatctgccta | agcaaaggca | tgcccttcta | acaagatttg | cttagagcag | aggtgataga | 180 |
| aggaagaatc | cgaagaccct | ctggcatggc | aatctgggag | cagcacattg | ttgatggagt | 240 |
| ccaagtgagc | acatttcaca | caattcattt | agtgacaagt | gggcttgctc | ccttttcatc | 300 |
| caggaaaaaa | actactcaca | gaccactgcc | cagaatctgg | aataagaacc | ctcattttaa | 360 |
| ggtattcttc | ccaacaaata | aatatctaaa | tattgaaaag | gggcatatca | gaaaacttaa | 420 |
| aagacacaat | aaccaaaaacc | aaaaccctct | tcaaaacaag | taagcaatgt | ctgtatttag | 480 |
| ttcactctaa | aacattctta | gcttttcttg | cagtttgttc | ctaaaagatt | tgattgggca | 540 |
| caagaggaac | gaaattatta | ataaaataaa | agcttatttt | tgtttttgct | gtggataatc | 600 |
| ggtacaaaac | gtttccagat | ctgagactta | aatggatctt | ttaaggtgaa | aaggagaatg | 660 |
| ccaggttcta | ctgaaat | | | | | 677 |

<210> 169

<211> 635

<212> DNA

<213> Homo sapien

<400> 169

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ttaagaagac | tgggcattta | tactctctct | tgctagtcag | cctggagcaa | gcttggagca | 60 |
| gacgcacatt | tttgtactgg | cacatattct | tagacgacca | attatagttt | atggagtaaa | 120 |
| atattacaag | agtttccggg | gagaaacttt | aggatatact | cggtttcaag | gtgtttatct | 180 |
| gcctttgttg | tgggaaacaga | gtttttgttg | gaaaagtcgg | attgctctgg | gttatacgag | 240 |
| gggccacttc | tctgcttttg | ttgccatgga | aaatgatggc | tatggcaacc | gaggtgctgg | 300 |
| tgctaatact | aataccgatg | atgatgtcac | catcacattt | ttgcctctgg | ttgacagtga | 360 |
| aaggaagcta | ctccatgtgc | acttcctttc | tgctcaggag | ctaggtaatg | aggaacagca | 420 |
| agaaaaactg | ctcagggagt | ggctggactg | ctgtgtgacg | gaggggggag | ttctggttgc | 480 |
| catgcagaaa | gagttctcgg | cgggcgaaat | cacccctggg | tcactcacat | ggtacaaaaa | 540 |
| tggttttgac | ccgctaccga | cagatccggc | cgggtacatc | cctgtctgat | ggagaggaag | 600 |
| atgaggatga | tgaagatgaa | tgaaaaaaa | aaaaa | | | 635 |

<210> 170

<211> 533

<212> DNA

<213> Homo sapien

<400> 170

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtgatctc | acaagtgtga | aaaatcttat | gaatgtaaaa | tgtgtggaga | ttcttctttg | 60 |
| tttttagctt | ccactttggg | aacatgtcaa | agcacacatt | gagaagtccc | atgagtgaaa | 120 |
| gagatgttgg | aaagcccttg | aacttggtcg | ttaggaaaca | tccacactga | agaggaacct | 180 |

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| gactgtatgg | aaggtcaaaa | aggctgtatt | aatttacatg | caaaaagtca | cactagagga | 240 |
| atgccatatc | agaatgcttt | tggtaaatat | acatgtttta | aagaggttat | atatcattaa | 300 |
| taaaaatatc | tagctgggtc | gaagaccctg | agttatctca | attgttcacg | gttacagatg | 360 |
| gaactcttta | ttattgagga | gttccactct | ttccccatt | tgctactact | acacttccct | 420 |
| agtctttaa | acaatttttag | gctgggtgca | gtggctcatt | cctgtaatcc | cagcactttg | 480 |
| aaaggccgaa | gcgagtggat | catttgaggt | caggagtctg | agaccagcct | gga | 533 |

<210> 171

<211> 568

<212> DNA

<213> Homo sapien

<400> 171

| | | | | | | | |
|------------|-----|------------|------------|------------|-------------|-------------|-----|
| cccttgsc | aa | actttccctt | aagtattgca | ctacaagtct | aagacacttt | tcactcaaag | 60 |
| ttccttcc | ct | ccttacctct | cttttaactt | ggagtcagac | tttcatcagt | ctgacaactt | 120 |
| ctccctgt | ct | ccttcccttt | cccccttca | caagcatttc | acctaacaaa | tttcttatgt | 180 |
| gcttaatcc | ct | cttagaag | cagatgccaa | gatgggatta | agcacataag | aggctcctgga | 240 |
| ctaataca | at | gacaaaggct | ccccttgaag | catcacacta | aaaggaaaaa | aaaaaaaaaa | 300 |
| acctagcca | tt | tacattaa | ctatttctaa | aatatagtat | ttgcttccct | atttgctaaa | 360 |
| acaaaatata | ct | aaacatga | ctattccaaa | aatctgtagg | gtactaagaa | tatgaagaga | 420 |
| ttcactctac | tt | caggggat | ggagttgtag | tagaaaaggc | tttgtggagg | gaggggtggtg | 480 |
| tttgaaatgt | act | tttaaaag | ccatcctcaa | agcctcgagg | gctataacctg | gcctggtgat | 540 |
| tatccaagga | cag | tccttc | aaacaggg | | | | 568 |

<210> 172

<211> 167

<212> DNA

<213> Homo sapien

<400> 172

| | | | | | | | |
|-----------|----|------------|------------|------------|------------|------------|-----|
| ccattttac | ag | gaatcagcca | cttcagttca | gacagcttta | ttaaaccgcc | tggagcgaat | 60 |
| tttogaag | ca | tgttttcctt | ccatacttgt | ccctgatgct | gaagaggaag | ttacttccct | 120 |
| gaggcact | tg | ctggaaacaa | gcactttgcc | aataaaaacg | agagagg | | 167 |

<210> 173

<211> 391

<212> DNA

<213> Homo sapien

<400> 173

| | | | | | | | |
|------------|-----|------------|-------------|------------|------------|------------|-----|
| cctcccaa | ag | tgctgggatt | acaggcatga | mccmccmcgc | cctgatgata | gacacgtttt | 60 |
| taacttcta | aa | aatatatga | tcattgattgt | gtctgtggag | acttgacat | atactaaatt | 120 |
| ttaamcaat | ag | agatatatt | gttcattacc | acattttggg | agtcattatt | tcctctatga | 180 |
| agagagaaa | ga | atttgata | caagttcaca | ggggcttcca | gtagattgag | acttttat | 240 |
| ctagctgag | c | tgctgatgta | tgaatttttt | ttgktattat | gactttcata | tgtattaaaa | 300 |
| ataaaatga | aaa | acaagg | attaggtgag | gaacctatac | gtctctaata | tgcaaaatac | 360 |
| cacagaaata | at | gactgktg | ggaaaattag | g | | | 391 |

<210> 174

<211> 474

<212> DNA

<213> Homo sapien

<400> 174

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| gaactcagag | agaggattgt | cacccttggc | atctgagctg | acactataag | gacaatgagg | 60 |
| agtctccttg | gggatatag | gggagatgga | aggacgatgc | ctgtcctacg | gggtcttggg | 120 |
| aggttaggga | tacacactgt | gagctgccac | aggctcaaca | gtacggatag | ggggtgctgg | 180 |
| aaccagccag | ggctctgatc | accaagctat | gtgccccatg | cagaggaagg | ggtagtggca | 240 |
| caactgaacca | cccagccaca | aggctatctc | cccatacagg | gcacctttaa | aaaaattatc | 300 |
| cttacagggg | aagacggggg | ggaaggatga | actgtgtgcg | gtgatgttgc | agtgagtgtg | 360 |
| agtttgtgtc | cgtccgcttg | tatgagggcc | taccttttac | taactagccc | ccaactttca | 420 |
| ttatctcccc | tttttctgtc | tacccttctg | ccttttttaa | gtggcttgca | atcc | 474 |

<210> 175

<211> 655

<212> DNA

<213> Homo sapien

<400> 175

| | | | | | | |
|-------------|------------|------------|-------------|-------------|-------------|-----|
| ccttgcaggg | gtggggatgt | gtgggcttgt | tcaactgttac | agcccatgta | tacctgaagg | 60 |
| gcaacatgta | cccacaaatg | ttccaggagg | taaataaaaa | atacaattca | gcctcttcta | 120 |
| aaccatcctt | gttgatatct | ctgctacttc | cgaaagttaa | ttcgttattt | ggactccata | 180 |
| atthtttcta | ttaattcacc | ctatgtccaa | ctccaacagt | gaaaaaaaatt | tatttaattct | 240 |
| ttgcaataag | cctataggca | ggcagcatta | tcctcagtct | gcagataaag | taaggctcag | 300 |
| agaagcttgt | atactgtcac | ttaggtagta | attgcaagag | ctggcattca | gaccagact | 360 |
| gtgggactcc | tcactccatt | ctctttcccc | ccactaggct | gctccttaaa | atacaatgga | 420 |
| tgtttgatga | acgcttggg | gaatcctggg | tggacacagt | tccttttctg | ccaaaagcac | 480 |
| cttgacgagt | tgtgaagaat | taatctggaa | aacttaacct | atthataaaa | acgtgttatt | 540 |
| aagggcaggt | tattcccacc | ccctttacca | aagaaacccg | ccctgacctt | tttttactgg | 600 |
| gggttgggtct | tgggcatttt | caacaagggg | ggaacagttt | aaaaattccc | ccctt | 655 |

<210> 176

<211> 660

<212> DNA

<213> Homo sapien

<400> 176

| | | | | | | |
|-------------|-------------|-------------|------------|-------------|-------------|-----|
| cctgggtcaaa | gtgggcatta | ccattcaagc | attactagac | atcacccgtaa | cgaaggctct | 60 |
| gttcacatga | aactaccctt | tctccattgg | gggctcagac | tctgctctca | tccaggatcc | 120 |
| tgaactctgc | tccaggcacc | tgttcaacct | tctctcccac | ccactgcctg | tcacttcaact | 180 |
| gactccagtt | acattgaaac | aattttcagt | ctaaggagg | atthttctacc | tttcagagct | 240 |
| gacctccgac | tttaagactt | gacagggtatt | tatcttgaaa | ccagagaggg | agctggagga | 300 |
| aaaaaaaaact | gagcaagcac | atcaatgcct | tttccaccct | tcttcatcct | ttccacactc | 360 |
| accgactgcc | attaccaaaa | cgccaagcac | aaccggtttg | gaacaagacg | cattccgttt | 420 |
| taattaaaac | caactcatta | tgtatthttag | tgggggggaa | gggggggcaca | atcagggttt | 480 |
| tcaccaccaa | atthttccaca | cggthttctga | acaccattgc | ctthtaaaaa | actatthttc | 540 |
| cacctccaaa | atthtttattt | aaatthttatt | tattacggag | gtgggtattct | tcctttggga | 600 |
| gccaaattgg | gaaatthtagg | gaacctthttt | tattaccggg | ttthttgggc | gggtaaaccc | 660 |

<210> 177

<211> 459

<212> DNA

<213> Homo sapien

<400> 177

| | | | | | | |
|-------------|------------|-------------|------------|-------------|------------|-----|
| ctthtttctct | tcctctgtgg | aatgggtgaaa | gagagatgcc | gtgkthttgaa | gagtaagatg | 60 |
| atgaaatgaw | thtttaattc | aagaamcatt | cagaamcata | ggaattaaaa | cttagagaaa | 120 |
| tgatctaatt | tcctgttca | cacaaacttt | actctthta | ctgatgattg | gatathttat | 180 |

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| tttagtgaaa | catcatcttg | ttagctaact | ttaaaaaatg | gatgtagaat | gattaaaggt | 240 |
| tggtatgatt | tttttttaat | gtatcagytt | gaacctagaa | tattgaatta | aaatgctgkc | 300 |
| tcagtatttt | aaaagcaaaa | aagggaatgg | aggaaaattg | catcttagac | catttttata | 360 |
| tgcaagtac | aatttgctgg | gctagaaatg | agataaagat | tatttatttt | tgktcatgyc | 420 |
| ttgkactttt | ctattaaaat | catttttacga | aaaaaaaaa | | | 459 |

<210> 178

<211> 720

<212> DNA

<213> Homo sapien

<400> 178

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ctgcaagctc | ccactccttc | catttatctt | aacgcccagg | ctgacttcta | agctgctttt | 60 |
| cactttccta | cctccactgc | attttcgccc | ctgataatth | ttgtaagctt | acctaagcct | 120 |
| cccttctttt | gagatccctt | tcttaaaagg | gtccattcta | ttaaccttac | cccatatcca | 180 |
| gttactttta | ctacctgctg | atctatcgct | accttggtcca | attcatggga | attacagggg | 240 |
| gcactgggac | aagagtaaaa | tgatccaaca | aacataatgt | tgcatthtaa | aaaataagct | 300 |
| aaaagatact | gatgactttt | tataactaca | acataatcgt | ttgtgaataa | gaacatatat | 360 |
| agtaaaaaga | tgaaaatgtg | aacaggttga | ctatttccta | aatttatggc | agaaggttgt | 420 |
| tctggagagg | atgggaagaa | aaaatgaagg | ctggcagtga | tgggtgggga | aatgcaacct | 480 |
| ccaaaattat | ctatctatat | atttttatta | aaaacaccca | cagtaattat | ggcaaatggt | 540 |
| aatggtttgt | ttgttctaag | gttttgata | catttaagat | ctcttgcttt | ctgggtacca | 600 |
| tttcttttct | tttcttttct | ttttttttca | aattaattcc | aaaagactta | tatctgctac | 660 |
| atgaagaacg | aagcaagttc | agctctcttg | gctgaaatgt | tcaaatgctt | gagggcaagg | 720 |

<210> 179

<211> 427

<212> DNA

<213> Homo sapien

<400> 179

| | | | | | | |
|-------------|------------|-------------|------------|------------|------------|-----|
| ctgtgaatct | gtctggttct | gaacttattt | tttagttatt | ggcaatcttt | gtattactat | 60 |
| ttcaatctct | tcctggttta | atctaggagg | gttgtatatt | tcagggaatt | tatccatctc | 120 |
| ttgtaagttt | tctagtttat | gcacataaac | gtgttcatag | tagccttgaa | taatcttttg | 180 |
| tattttctgtg | atatcagttg | taatatctcc | catttcattt | ctaattgagc | ttatttgaaa | 240 |
| cttctctctt | cttggttaat | cttgctaattg | gtctatcagt | tttatttatc | ttttcaaaga | 300 |
| accagctttt | tgtttcattt | atcttttgta | ttgtttttgt | ttgtctcaat | ttcatttagt | 360 |
| tctgctctga | tcttcgttat | ttcttttctt | ctcctggggt | tgggtttaga | ttgttcttgg | 420 |
| tttctct | | | | | | 427 |

<210> 180

<211> 728

<212> DNA

<213> Homo sapien

<400> 180

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|-----|
| caaacacaaa | agtcactgtg | tgtgtgatgc | ttctccaatt | ccactcatcc | tggctgccat | 60 |
| tcatgcacta | gtgcatgtat | gcattttttac | attttttaaa | ttacaaaaat | caacctatta | 120 |
| taactgctta | gatatatatg | aagtaaaaat | gaaagttctc | cctttacatg | acctatcccc | 180 |
| catcatttcc | ctcttttatct | tatactgtca | gcattcccag | cttgtagcac | agtgtctggc | 240 |
| aatagtaaat | octcaaaaaa | tgatcaatga | ataatttaat | aatgattaat | aaataaatta | 300 |
| atgatgatgg | tgaagataaa | ttttagcatt | tattgaacgc | taactacaaa | ccaggggagt | 360 |
| tggtaaatat | tttataaaaa | tcaatgaatg | agctaaaatg | ccattctatt | atttttttgg | 420 |
| atacggttta | atatttttact | cataaatatg | cttaaagaat | attataatta | tatgacttag | 480 |

<220>
 <221> misc_feature
 <222> (1)...(700)
 <223> n = A,T,C or G

<400> 184
 ccaggscawt gaggaaaaagr gaaagaatwt arrggstwt caaataggaa aaraggaagt 60
 ccaaattggt ccntgttkg ccagataacc atgattgk gk atttagaaam ccccatgwt 120
 tcagcccaaa atctccttaa gctgattaag camcttcagt aaaktctcag gataaaaaat 180
 caatgtgcaa aawtcacaag crttcctatm cgamcaatam cagmcaaaca gagccaawtc 240
 atgagtgrac tcttattcac aattgctagt aagagaagaa aatmcctagg aatacaactt 300
 mcaagggatg tgaaggwtct cttcaaagaa gaactacaar ccrctgctca aggaaataag 360
 agaggmcmca agtaaatggg aaaagcattc tatgctcatg gataggaaga atcaatcccg 420
 tgaaaatggk gatactgcc aaaataattt atagattcaa tgctatcccc atcaagctac 480
 cattgacttt cttcmcgaa ttnggaaaaa tctactttac acttyatagg gracaaaaa 540
 agaagcccw ttagccaaga caatcctagg caaaaaagac caamcctgga ggcatcacag 600
 tmcytgactt cmaactatwc taccaaggny tmcrgkgmcc aaaacagcac ggkacntggt 660
 mccaaccrg acwtwtwgac cmmcagacac agaacmgagg 700

<210> 185
 <211> 192
 <212> DNA
 <213> Homo sapien

<400> 185
 ccagyctttt ttttaagtaa ggcgtttttt aagctcattg tagctacaaa gtcaataaat 60
 tgggtctttgt tattttttacc tgaaaaggct gttaaagggt aaaatgacaa actcaaattc 120
 aaagggattg gaggatttgg tgtttatgat ttctcagaac aacaatctag agaccaccag 180
 ggtgggtttc ag 192

<210> 186
 <211> 688
 <212> DNA
 <213> Homo sapien

<400> 186
 gtgctggaat tcgcccttag cgtggctcgc gccgagggtg gatattttct ctggatagat 60
 ttcagatagg tagttccctc aaataagatt atatgggttt gcattttcaa ggcagagttg 120
 tatacttctt gctctttatt taaataaaaa aacttgaaaa tctgttctgc ccagtattgt 180
 aagcgctcag gtacaaatat gaatgaaaca atctctgcct aagtaacaca agtataggga 240
 caagattctc agtaaaattc tcacgtgaaa tttgtaactc actagacact atcaggagat 300
 caataattat gtaattaaaa aaaataatta cctgccaaac tgggtttctt tttggcactt 360
 ctgcttggtt ttaagacaat tctcacatag aagcttatta ttccccatta gtcattccat 420
 agatgtaaaa ctggtagaaa caggacttga attgaacatt ctttacaagt aagttatata 480
 gcttctgaaa aaagggcttg aaaaagcatt tttggggact ataagaacct tcaaatgctt 540
 tcccccttta acaaacctta aaattatatt gaaaataatt taagggggct gattttctct 600
 tgtaaaaatc ttgaacccca cttaccaggt ggttggtcaa accaaagttc aaaaaaaagc 660
 ttctggcctt tcttttatcc cacttgca 688

<210> 187
 <211> 779
 <212> DNA
 <213> Homo sapien

<400> 187

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| gcaaaaaaca | gatacatttt | cagtgtttta | aaatgaacaa | gtatggaaag | gcttatacag | 60 |
| taactgaaaa | gtctcctttg | ggaagccaag | gtgggaggat | tgcttgagg | caggagttca | 120 |
| agaccagccc | aagcaacatg | gcgagacccc | atctctacaa | aaaattaaaa | aatcagccag | 180 |
| gcatggcgga | catacttgta | gtagtaacta | catgggaggc | tgaggcgga | ggatcacttg | 240 |
| agtccgagag | tttgaggctg | cagtgagccg | caacgcgccc | tgtactccag | cctgggcaac | 300 |
| agagcaagat | gctgctctaa | aagaaatttt | cttttaaaga | aaaaagtctc | cctcatagcc | 360 |
| tgttctacaa | aagtcctatt | tcttcccaca | aaaagcctct | ggtacctggt | gttagttctt | 420 |
| ggggtggaag | attactttta | aaaatagaac | tattttttta | gtatatcttt | tagggaactt | 480 |
| tagttcccga | agcttttaga | aatgggatct | tgaaaaacaa | agggatttca | atacctatga | 540 |
| caatgcttaa | agaattattg | gggcatttat | ttttcaatgg | aggggtccaca | aatctttgga | 600 |
| aacccttggc | caattaccag | aagccacttt | aatttttgac | cgaaaaatgtt | tttaaaaaatt | 660 |
| ggcttttggg | aaaactgtct | ctttcccca | aaatgaaaac | cttgaaaaaa | aggggaattt | 720 |
| ttaaggttgc | cccctcatta | aattttaacc | cctctgaaag | aaaaccctct | tgtgacagg | 779 |

<210> 188

<211> 394

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (394)

<223> n = A,T,C or G

<400> 188

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ggcgamgtct | ggyccaccatc | atgcccttta | atcaactcac | acctgtttta | agagtgtttc | 60 |
| tgatttgacc | ttcatccctt | agtttactgg | cgtaaaaaaa | agtctcagca | attttcatta | 120 |
| tttctcgtgg | gtctcattat | caaaccctta | cttatttcgg | catatttcct | ctgggcttct | 180 |
| tctagtttct | gccttacaag | caatgctgtt | ctgtaaattt | attgaaacct | ctggaacatt | 240 |
| tcacctttag | agatggagga | tggaaggatt | ggyaccagaa | gagggctaag | atacgtttct | 300 |
| tgtcttngag | ctgaaagcac | agycactctt | ccttcgtttt | gycgatgaga | aaagttgagg | 360 |
| ccagaaggga | ggtgacatgt | ttagagtcac | ccag | | | 394 |

<210> 189

<211> 681

<212> DNA

<213> Homo sapien

<400> 189

| | | | | | | |
|------------|-------------|-------------|------------|------------|-------------|-----|
| aagttctgac | tttggctctat | aaaacagggt | tattggctgt | ggctgcactc | aatatctaaa | 60 |
| aagttattag | gaagtgcctc | gttattgtca | ttaaagatat | ctaaatatgg | tagaccaaag | 120 |
| gttgttgaga | aacacatatt | atggactgag | ttctgtttct | tctgctgtgg | cgcacctaa | 180 |
| ctcaagcctt | ccttctctcc | ctccccctct | ggccggcatg | gtatctgagc | tcacagacag | 240 |
| acaaggcatg | ttagaatcat | cagatcatga | gcaccgtgct | gggatttagc | cctctccaaa | 300 |
| gtcaattctt | acagtccata | ctttgcttaa | atcctcagtt | gttgaggctc | gctctgctgt | 360 |
| cagtaatccc | agctataaat | ttcccccaaa | tgtggggcct | agataaagta | gaagggtggat | 420 |
| ggactcagct | tattttcatg | ggatgacagg | aactggaaag | agaaagggca | ttgaaaataa | 480 |
| aaagttattc | cagaatagca | ttaacctctt | tactgttcaa | gaattaagaa | agcctactta | 540 |
| gaaatgaggg | ccttgagaat | gatacccaaa | tattggctct | tctacaaaaa | aatggccttt | 600 |
| ccaaatatct | gctttcctgt | tcccccaattg | gctttttaag | tagaattaag | ttacctaaaa | 660 |
| ctttacctga | aggggtggtt | t | | | | 681 |

<210> 190
 <211> 839
 <212> DNA
 <213> Homo sapien

<400> 190
 caaatacatg atttccattg gcatagactc ttctatagtc tctcaggcac accttatgac 60
 taataagaac actgtcttct agatataagc caagtttttag gagttatctt tgtagtttct 120
 gtgttgagac tatgggtctt ccctgtgcaa agacttgatt agcaaatact atttgaaacg 180
 atcccaaatt catagtgcag ttgaccaccc ttctgatcaa ggggatctct gtatatccca 240
 tgaaagcttc ataggtctca ccctagatta agtgcttcac ttctcaagac agtgaacaga 300
 tggaagactt ttgtagttat cattatacaa ctgtgccctg tgtgttttat tataacaacca 360
 gagaactgag gcactggctt tacctgtcag ctacgccagg ggtgtgacgt catctttctg 420
 acttgatcac acatgccaca ttgcttaata tttcaagctt agactgaaat aatcctgtgg 480
 taaaaaattt ttggggggct ggggaggtaa agaacaaggg ggggaacttt ggaatatattt 540
 tattcattaa tcatatttcc cgaattgtat tttattttga aatgaccata agggacttaa 600
 atacgtattg tggttaaatt aaatggaccc aaatggaggt aagtaaacct aatgggacaa 660
 atgaataaaa ggtttatgac tgggagcatt taccatgaa cctccttaga agctatttaa 720
 cctttctttt ggaaagccct gaaggctggg aacttaaatt ttaaagacag tacctatttc 780
 cagaatcgct tccaaatggc catgttttaa agggccaaca ttttgggatg gccctgccc 839

<210> 191
 <211> 697
 <212> DNA
 <213> Homo sapien

<400> 191
 ccatactgaa tactgatttt ctaatggaac tctattcaat ggcgattgta aaacctgag 60
 gctccgttac tattatggag catactttca tctcattctc ggctattggg caatatgtat 120
 ctcataagat tttatcacat ttcacagatg aactgttaat tgattccatg ggtacgatta 180
 ggcgagatcc aagctggagc tgcagctctg agtcccataa attctttgtg cttctgtaaa 240
 gaataaatct gtttttaatg caaattaaaa ctactggcag ggaatttttg cttccagtta 300
 ttaaaagact ggaaatgtgt aagtggagaa aggcaataac tgcagtaatc tcttaccgga 360
 ctctattata attccaaaca tacataatgg tgagaaaaac cgggaaggga agaattgtggc 420
 aatgtccact ctttgcccca aacataaccc ttaattttcca tggcggggccc aaacactggg 480
 aaaaacccaa atggtaccct ctatagcatg caacttttat ttcactccaa acgaaaaatt 540
 attttgacta tggtctggga aatccattag tagaagaagt tttataacct ataggaaccc 600
 ggccatttca tttctaccaa atcacaggaa ttttagaatg ggcaaggaat ttacaggaag 660
 acttgcccaa ttatcttttt ttgggggact aaaccaa 697

<210> 192
 <211> 687
 <212> DNA
 <213> Homo sapien

<400> 192
 ctggttacta tagctttgta gtataattta aagtcaggta atgtgattct tccagttttg 60
 ttattttctgc ttaggatagc tttggctatt ctggatcggt tgtggttcca tataaatttt 120
 aggatagttt tttgctattt ctgtgaagag tgtcattggg actttgatag ggattgcatt 180
 gaatctgaag attgcttttg gtagtatgaa cattttaaca atattgattc ttccgattaa 240
 tgaacatgga atgtttttcc tttatttggc gctctcttta atttccttca tcagtggttt 300
 ataggtttca ttatagagat ctttccttct tttgggtaat tcctacgtat ttaatttatg 360
 tatcgctatt gctaaatgga atgacttttt aaatttcttt ttcacattgc tcttggtggc 420
 atattaaaag ctactgatgg atggtgattt tggattctgc cactttactg gaattgggtg 480

tatttggttt aatgggtttg tctaattctc ttcaatcaat aaaattgtgc gtattttaact 180
 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 229

<210> 196
 <211> 557
 <212> DNA
 <213> Homo sapien

<400> 196
 gcggtggctc atgcctgtaa tcccaccact ttgggaggct gaggtgggca gatcacttca 60
 agttgagagt ttgagaccag cctgggcaac ataacaaagt gagatcttat ctctacaaaa 120
 aaattaaaca aacaaaaaaaa caaatcaaca ttcatttgca gggctctttg gtcttcttaa 180
 agaacaaaca tatgaaataa ataagctgat tcttaaagat aacaaatata atgagctttc 240
 tcaactgtaa aagcatctct aagttgttct atcaatgcat atccactcca tgaactaacc 300
 tgaagaaagt gttgaccatt ctaccaatt aactgtaaac taagattgct ttaatgggtt 360
 gcctaaattt gagtaccttt aaatttttgc tttttatcca aattcattct cccttcttca 420
 aattaaatag ttttgtaga aatcggataa gcaagatgta ctttttagaa agggcaatag 480
 aatcctacaa catgctagaa tttgaaatgt ttttttaaata cagtmmtttc tctatgctag 540
 taactaagaa aattata 557

<210> 197
 <211> 624
 <212> DNA
 <213> Homo sapien

<400> 197
 ttttactacc tatatttaaa atgatccctg acgcccctca agacaaatat attaattttt 60
 ttactttgtg ggatagagat cagaaaaaga gtagagatga aaatactgga gaaacaatgc 120
 aggagatatt tatgaggtga gaatgtcaag aaacttgtaa agggagaata ctataatgac 180
 ccctgaagag agagcttttag accagttgag tattagaggt tgccacgtgg ctattcatcc 240
 actaataaat acaagaaatt actaaaatgg aagccactgg aaatatgttt tgaggaagggt 300
 gagaatgtgg acctattata aatgggtgaa tatgatttct ttctcattaa gttcataaat 360
 aactttcaga catgtaacag tttatgaagt gtgccgtagt catttagtat aagtttttata 420
 cacaaaagtg tttttactaa gactgtcaca ggttcttttg tgaatcttgt ttgtttttcc 480
 tcattgtaaa tactgcaata gaacatttgt gtcttaacat aaggcaataa atgaccttaa 540
 gaaccttcac ttttatatag aaagtggagg aaaagttggc agagtaattt gttgattata 600
 gataaaagct cttgtagaaa ttgg 624

<210> 198
 <211> 175
 <212> DNA
 <213> Homo sapien

<400> 198
 tttttttttt tttttttttt ctaacactta tgcatttatt ttcattgtgta agaagaaaaa 60
 cgtaactagc acgtgaacat gactgcatgg atacacggct cagcacgagg ctaaagtcag 120
 aagtgagtga aagcaaaacc gcatgttgat ttaagtgaata taacagaaca gaaaa 175

<210> 199
 <211> 871
 <212> DNA
 <213> Homo sapien

<400> 199

```

ctgttgatca atgatgagct cccaagagta accagcctct atagatcag catcactggt      60
ttctcaggaa aagcatcacc attgttcac ttgctgcaaa atgtatgcac aagtatcttt      120
ttatttttaa aaaagccctg acattttatg actgctgctt ttctaagata ttttcaaata      180
tacagtccat acgggttcaga cacaatggac tggggataga gacggctata gtgccgataa      240
tggagaaaact agccagagct tcagatattt gttttccagg acatctcaat aattgggtac      300
acctcacaat atgtgagact tgacgtcgag tggcacggca tactctggcg caggcacttg      360
ataaagactg tgtttgcaaa tacttagcct gcacttcaag ataccaggca tctaagcacg      420
tcccagatgg tgacagttaa tcttcaaaaa accctatgtg gaagtattat cattgtcctc      480
attttacaga tgaggaaaaa gagacacagg gatgtcaata tcttcctcaa ggtcacacag      540
caagtaagtg atggaacagt ggctcagcca tgaagctatt gctgttaacc actaggttga      600
tttgccctta ttaatttctt cctaaaactg cacatttccc gttagtccct ctttttggtc      660
tgtcgtttga ctcttggtta ctgcttagag gaagattcat tctattattt tctaacttag      720
taaatatgtg caactccctg gggacatgac caggcaaaag ctggatacag aaatgtatgc      780
ccaaacacca tcccaagtta cccctaacag gtcttttctg gacctgttt gtaagggggg      840
tatatttgga aaaattttta aaattttctg g

```

```

<210> 200
<211> 737
<212> DNA
<213> Homo sapien

```

```

<400> 200
gacattttga aggtaacagc aatatctgtg tatagatggg gttgtggttt tgttatttat      60
ctgctattgc tgaactatcc tttgtcttga gcgataaaaag agaagtaaaa tactaaagaa      120
ctgaactgtc catttctgga ccatgagtaa agatgctggc tgtcaaactt cctgttcata      180
cattagttta tttatagagt gtactctcta tgtaaggat tgactgataa tgttactttg      240
acttcagata gcttgagtt taatggagga agaagacaaa catgcaaata actagggtcaa      300
tgaggcatcc tttgtgttcc attggaagct aggtctgctt gtaaccttgt taatttctgt      360
ggttttggag tgcattcatt agcaaataca cccctgttcc ttatccattc tctgcttttt      420
tctttatttg gcatttgatg acattttttc atgtggggaa attgagtcag gtgaggtgga      480
aagaaaataa ggacacgaca ctaaattctt tgatgttttt ccttaaaaaa ttgtttttca      540
agtgtcccat aaagggttgt gaagttttta gagccatagg acttggatta ttgtgaaaga      600
gtgtctctag ggggccagggt taaaccattt caaggactct ccttctctca tctcccttgt      660
tccaccaggg gtggcgaccc ccaaaaagca caaagcctcc ctttcttcat gggaagggta      720
aggaacggaa gggaacc

```

```

<210> 201
<211> 493
<212> DNA
<213> Homo sapien

```

```

<400> 201
tctagaaatg cagctttttat ttattacccc atttctttca agtccttgga aaataacata      60
ttaagggtac aagaaattaa cacatgatgg aaaagtcatt gtgacgccaa tgaatttcat      120
tgagtataaa ctcatctact tcaaatttat tttataacac aacctaatg actcaagata      180
attatttaat ggttagctct taagttgaat tgggtctacat aatgcgtggg aagaaaacca      240
gatttttagc cttcttgcca aatccagacc tctggttgat ttttctttga cagaagatgc      300
aagttatttt ccaatttcac aattaaatgt atttaacatg aacattattt tgcttttaaa      360
actataaaca ttgtaggaga attatagcca gtcttcagtt ataaccactc caccctcctc      420
actttctctc tctctctctc tttttttttt gctatgggat ttaatgggaa aaatatgtaa      480
aaactgtcac taa

```

```

<210> 202
<211> 283

```

<212> DNA
<213> Homo sapien

<400> 202
cctttttatc tcagtgcacac cgtccgggga cgcaggtggt ggtgactcaa ggctagcctc 60
aaagggcagc cccacctcct catcctggac cacagagacc acctgcttgg cgcgccgtcg 120
cttttccgag aggggtggctg actccgggggt gctggggctg gggctgccgc ccccgccgct 180
gttgctgtac tcctcgcccc agtcgatggg ggctgccctc ggacagcagg tgcaggttgg 240
gggcactggt acgcaagacc atgctgcccc gagaggtaga tct 283

<210> 203
<211> 713
<212> DNA
<213> Homo sapien

<400> 203
ctgcttttgc gcaaggtgcc actggacgag cgcctcgtct tctcggggaa cctcttccag 60
caccaggagg acagcaagaa gtggagaaac cgttccagcc tcgtgcccc caactacggg 120
ctggtgctct acgaaaacaa agcggcctat gagcggcagg tcccaccacg agccgtcatc 180
aacagtgcag gctacaaaat cctcacgtcc gtggaccaat acctggagct cattggcaac 240
tccttaccag ggaccacggc aaagtcgggc agtgccccca tctcaagtgc cccacacag 300
ttcccgctca tcctctggca tccttatgcy cgtcactact acttctgcat gatgacagaa 360
gccgagcagg acaagtggca ggctgtgctg caggactgca tccggcactg caacaatgga 420
atccctgagg actccaaggt agagggccct gcgttcacag atgccatccg catgtaccga 480
cagtccaagg agctgtacgg cacttgggag atgctgtgtg ggaacgagggt gcagatcctg 540
agcaacctgg tgatggagga gctgggccct gagctgaagg cagagctcgg cccgcggctg 600
aaggggaaac ccgcaggagc ggcaccgcag gtggatccag atcttcggac gccgtgtacc 660
acatggtgta cgagcaggcc aaaggcgcgc cttcgaagga gggggctgtc caa 713

<210> 204
<211> 275
<212> DNA
<213> Homo sapien

<400> 204
gtagacaagt acagcagatc cagacaccag atctagctag gctaaatgta cagtatctaa 60
cttgatctga actgaacctg tattccttga tgatgcctaa aactacatcc atagaattct 120
ggatgaacctg taatacagtt ctgaaagtac agttttatat aataagatgc tgatctcttt 180
attctttcaa gtaagagtgc tagagaacaa attgtgttac ttgccttggg atttattgaa 240
cgtctggaaa atgctgtctt cctagatcca aacag 275

<210> 205
<211> 694
<212> DNA
<213> Homo sapien

<400> 205
ctgttcctgt acattttaact gaaaaaaaaaag taacttaaaa taatataaaa atagcactca 60
tgtatgtcct acagttatag gtgaaatttg atattgtttg tcttacatag catacctata 120
gacagcttaa gtaaagtgc tgttaagagg gttatgctta ttgatgaact cttgtagtgtg 180
cttaccagct ctgttagtat agttaaattg atctcagtag cttcaagtat ttataaaatg 240
gttgaagtcc aaatacatgt gataattaca atacactttg aattaatgga ggggtgggagg 300
ctagttgaaa tgcattttat ttacccaagg agtatgttaa aatgatagtt ataaatgttg 360
gaagtttaaa gcaagatact cagtttagtt ctttacaaat cataagaaga acaaaattag 420


```

atgttgacat tgctatttta ggctgtgtgt tttccatatg cttcttgctt tccctgtcac 480
aggtgggtggc agcaatattg gtgtgattga ggttatgctg gcaccactcg cacacaggcg 540
cacaatgggtg ttagctgggc agaaagagtg gcattctctg ctaccgggct gggggcgacc 600
tttaccatag gatgaagtaa ccttgcattc ggctgcaagg tgtactgtac cgtacacagg 660
tgctgggtcg atggccactt tctgcttttc tttc 694

```

```

<210> 206
<211> 704
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(704)
<223> n = A,T,C or G

```

```

<400> 206
tttttttttg gnaaaaacag ggtttcatca tgtttgccag gctagtctca aactgctgac 60
ctcaggggat ttgcccgcct caccacaattc aactttcgta agtcagtatt taccatctaa 120
ctcagtgctc caaaatttaa aatttccttg cactttacag caaaaatata tattggggct 180
ctactgaagc aatatatata tgtcaaaact aaaaatcaga aaagcaaaag ggtccattca 240
acatatagca gcttatattt aaatatgtac aggtatgtat gttttcacag ttagatcttt 300
aaaaaaattt atatttgata tgttcaaaaa tacttctatt ggctataaat aatattttaa 360
aagctcaact gatcaaaatg cattccaaga acatatcaaa ttaaataaat cttctacgtc 420
tttaaaaaca gataattgaa gtcagtaaag cttgaggttt gtgttaagtg tattctgtca 480
gtccctacta ctagggaagg cagaatcttc taaatacgat acgaaagaaa ctcccaaagc 540
ttggaaggaa tcggcagctc ctgaactttt tggggggggc atccctcttc gggattgaca 600
tgcgacataa atgttgcaag ctaagggacc cccccggggg gagtgggccc caaaaaaac 660
cacaccttcc ccgtcaatgg tggccccccc accaacctta aaaa 704

```

```

<210> 207
<211> 225
<212> DNA
<213> Homo sapien

```

```

<400> 207
ccattttaac tgtactgcc aatagaattct ggaattgtgg aaaattgtat cattgaagtt 60
cagtaggatg tgtggcttaa aaatttatca ggaccacaaa aaagaaaaca aaaatatttg 120
gtactgaggt tcattgccag ggcaggaggt atttccagaa aatactcatg cctgtgttct 180
gttccttgct ttcccaaata ctgcatgtga ctttcctaag cggca 225

```

```

<210> 208
<211> 678
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(678)
<223> n = A,T,C or G

```

```

<400> 208
cctatatcta tcaaaaaaaaa tccagttcct aactaataat ctccccaaaaa gaaagcacca 60
ggaccagatg atataaatgg caaatttttt caatcattta aggacaaaat aataccaatt 120

```

```

ctgtatcatt tcttccagaa cacttccata ctcacgtat gaggccagca tcactctaatt 180
agcaaaacca gataaagcca ttacaagaga gagtgacaga ccaatgtggt tttattgagg 240
atgcaaacaa aatttaacat aatatttaaat agtgaaaaac tggatgctct ttccttaagt 300
tagagattaa ggaaagaatg tccccctcac tactcccata caacacctta ctgaaaattc 360
tagctagctt tataaaataa anaaaaacca naaaataaaa taaaagggtg acagactgga 420
agatacagtg aaggaggaag aaataaaatt ttctttgctc ataacatgat tcttctatgt 480
ggaaatcaca gagatttgaa catttttttt ttttgagaca gtttttgctc ttgttgccca 540
ggttggagtg taatggcgcg atctcggtc actgcaacct tcacctcccg aattcaaggt 600
gattctcctg ccctcagcct tcccgagta agcttgggga ttaacagggc atggcacccc 660
ccatgcccc agctaaat 678

```

```

<210> 209
<211> 720
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(720)
<223> n = A,T,C or G

```

```

<400> 209
attattttga accctagcat ttagaatga aaaacttttt ataacaatca aatacatgat 60
aaagtatgca aagagtagga aattattctg atgacatatg gagggttaca aaggagaaaa 120
ctttttgcta cctctgataa agaatagact aaattctcca agaccaatct gactggtgtc 180
ataataaaaag gaggtacaca cggaagcaca agggatgtgt gcctctggag gaaaggtcag 240
gtgaggactc agtgagaaga caagccaagg agccaggtct tggaagaagt caaccctgtt 300
gacaccttga tcttgacta accctgtgga caccttgatc ttggactttt agcttccaga 360
actgcnagaa aataaaattt tcttgtttaa gccaccana gtgtantgtt ttgttatggc 420
agccctaaca aattaaaatt atattttaac agagaatata aaattctaata ataacatttt 480
acagtaaagc attcatggtc ttttttttct tattaataaa tccatcaaaa cagaaagttt 540
tgcaaaattt taacacattt ctctaccact actgtttcta ctctcttaaa actactccgc 600
aaatataaaa atagaaggcc aaaatgcac attaaaacga tgtttgggga ctaatggcct 660
taaaattcta ttacacttgg aaatatacaa atattcaaag attatctatt gatcacctca 720

```

```

<210> 210
<211> 277
<212> DNA
<213> Homo sapien

```

```

<400> 210
tccatgtatt tttatacaga atggaacaat atgtatgtat gcaatyktta cattccacca 60
tgaaataaaa cagtataatg aaaataacaa tagattcaaa caatgatatg ctattttttt 120
ttacctatga cattggcaag gtcttcttaa aaaatctgcg aataaccgat gttggagaga 180
tcatggggaa atagccactc aaatgttact catgagagtg tacatatgtg taacttcact 240
tgaggggcaa tttggtgata catttaaaaa gtttttgg 277

```

```

<210> 211
<211> 715
<212> DNA
<213> Homo sapien

```

```

<400> 211
gtggtagaaa tactaatttt gcaattacag aaaaaaacia atgccattca catgggttyct 60

```

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| aacaaaaagt | gtctgaccac | ccccacccc | caccctcaa | aaagccctta | aataaagagg | 120 |
| aagatcaaaa | gaaaacaaaa | taattcccga | gtttcacctc | atacatacaa | tatagcacag | 180 |
| gaagtggcaa | agtttaaaat | aatgccttta | ctgttaggac | tagtatgctg | tcaaaagcca | 240 |
| caatcctttt | gttttagtga | gttgattttc | aatagaaaaa | tacaaatgaa | catgtgttta | 300 |
| agttccaaca | tggattgagc | acctctgaat | ttagtatcaa | atgattaatt | ttatttttca | 360 |
| gatgtcaaat | cttagtataa | aattttccat | tattttaaac | ttcacttgaa | tctttaaaaa | 420 |
| agctgtctaa | attgtactat | atgagttcag | tttaatcttc | tgtaaaatgc | taacaaattg | 480 |
| aactgtcagc | agtcttttaa | aaaaaaatgg | gggctgggtt | atttctagaa | gaactctcat | 540 |
| taagctttga | aaatcagaaa | tcagagacaa | ataacttcag | atatagacta | gctccacaag | 600 |
| caaattttata | caattatctg | taacagtcta | tacatatatg | tgtatatata | tataccgtaa | 660 |
| ccactttcat | aggtaaaaaa | tattaacttc | atgtcacact | atgatcagaa | gtata | 715 |

<210> 212

<211> 717

<212> DNA

<213> Homo sapien

<400> 212

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| agcctccccc | aatgccttaa | aaggtcacag | tagatctcag | ctctgaacag | aaactcaact | 60 |
| gaaactcttc | ccacaaccca | gcagtagata | tattaaaacc | tacaattttc | agggatacaa | 120 |
| ccaatattta | attcttttga | gggttttggt | tttaatacaa | ggacacaaac | acacgtataa | 180 |
| aatgacgatg | tcaatactga | ttaaacagaa | caacaaaata | agaagctcaa | attatcatca | 240 |
| gctattgtgt | atatctgaaa | taacaataat | gcacttgatt | ctgaaagaat | gatttagagt | 300 |
| cctactctga | aaatctaatt | gtcttgatgt | ggcgaagtga | gaagaaagga | tgatttttct | 360 |
| aatgaaaagc | atgtatacgg | gtagcccttt | gcgagattct | gtcaaaaccc | tgaattttgc | 420 |
| attagcttgt | ttaccaccca | aacgttttta | cccgaggatg | tgcagcaatg | ggaactctca | 480 |
| tacactgctt | gtgggaatat | aaatcagtat | aaccactttg | gaaaaccatt | taacattgtc | 540 |
| aactacagct | ctacacacaa | gtgctataac | caccatttcc | actccagggg | atacaccta | 600 |
| aaaatatgaa | gtgcccattg | ctacccaaaa | ggccgcctaa | aaggaatgct | tttgagaagg | 660 |
| gttaaccttg | ttaattagtg | gcaaaactgg | gaaaacaacc | cccaaagggt | cccatcc | 717 |

<210> 213

<211> 599

<212> DNA

<213> Homo sapien

<400> 213

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| cctgtttttg | cgaggcagga | gggaagcggg | atgggagtg | tggttaggcc | aagggtagtt | 60 |
| caaagcgatt | cagcaggatg | atgaccacag | gagtgtcgtg | gccgggcctt | tcagcccccg | 120 |
| tgtggatgat | gaccggccat | ccaggacatg | cgagggtctg | ggacagtgga | cagccagtgc | 180 |
| cacacaagga | aggaccgatt | aaatgacaca | gttaaaggaa | tttggcctag | ggagtgcag | 240 |
| ccagaaaagg | ttggcttttt | tatatatgta | acattggaaa | aaaggaacat | ctcctgttcc | 300 |
| ctgtattaag | ttttgacttt | agctcagcaa | atgcagtgtt | tgtggcagta | aatatactct | 360 |
| gataacaatg | ttctttccca | ggaatttaga | gttttatgat | ggttattgaa | aatgtttaca | 420 |
| tgacaggctg | tcaataatat | tttttgcttc | taaaaataaa | acatacataa | agtgtacgga | 480 |
| ttttaagtat | gcaactcact | gaacttttca | taccgtaata | caccacccta | gtaaccctcc | 540 |
| cccagttcaa | gatgtagact | gtttccaata | acccctcatc | ctgttccctta | atagcccccc | 599 |

<210> 214

<211> 789

<212> DNA

<213> Homo sapien

<400> 214

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ccttatgaca | aaccttgcta | tgccaaggat | atgcttcact | atcttcatct | atcaaaacac | 60 |
| tatgcatcat | agatatctaa | ttttttcacc | tcttgcatga | agtctttcct | gatttccctc | 120 |
| tgctgaaatt | tctctcttca | aatgatgtgt | ttccatagta | ctttgtccct | tttcaaagat | 180 |
| atatctcaca | tcgcatatct | taccacagtt | agtttcattt | cttaactctc | acactagatt | 240 |
| acaaagtcaa | tatagacaaa | gaaatgttca | accttatata | acctcctctg | cctatgctgg | 300 |
| taaattgcac | ctactatgtg | ttcaataaga | gcttgtcttt | ttcaatatac | aaaactttgt | 360 |
| aaagattaaa | gaccttgtag | aaagtcaaga | ggaagatagc | aatttcactt | ctaagaactt | 420 |
| accctaagga | aacattcatg | aagagataca | aggggttatg | tgcatggatg | ttcattatca | 480 |
| tattattctt | cattatgaag | attatgatgg | taataatgaa | aatgattatc | ttgtattggg | 540 |
| ccttatttga | agtcaagcat | tgagaatgta | ctttatctgc | attatctcac | tgagttctcg | 600 |
| tagcagccct | ataagggtaca | gactgttatc | taagcttaaa | aaaataaagt | taatgtccaa | 660 |
| ggtcaaacaa | ctagtaaaaag | aagggggcta | ggaaatttgg | aaccccaaaa | ggggcaacct | 720 |
| ctcaagggct | atgaatcctt | accattatta | taaggaagct | tggcccatgg | tggcccaaaa | 780 |
| aaaaccggg | | | | | | 789 |

<210> 215

<211> 765

<212> DNA

<213> Homo sapien

<400> 215

| | | | | | | |
|------------|-------------|------------|-------------|------------|------------|-----|
| ggatgtctga | gcaggagaga | gaccatgtga | aggatggact | gaatggagac | ttgtatcaaa | 60 |
| gagtcctgag | atcaaagact | tgtattagag | aggggtgttg | tagtaatcta | gtcagggtat | 120 |
| gagaaatggg | ttgtattaga | gtgtcaggag | tagtcgtggc | aaaaatatat | agatcaggat | 180 |
| gagggatggg | cctcatctca | caccctgact | ccagtcaatg | gcagtggctc | cctggagtag | 240 |
| actactatag | gaaggatttt | gtaaaagttt | gtctggcctc | agtggagggt | gaggtagggg | 300 |
| aggagttcta | tgaacagtta | gtggtgtctg | ccatggttga | aacaatggag | aagggggaca | 360 |
| ccttttctgt | gcagatgttg | cttctggtag | atataatcca | caatgtaatg | ggagaagtac | 420 |
| taagaatcag | taaattatgg | aggggtgtaa | agactactga | tatttaagcc | tgcggaaccg | 480 |
| acttagagaa | atgatagtta | aaggagaaat | atccagcaaa | caaagatatg | acattgaagt | 540 |
| ttgggactgc | gatttagtacc | agagatttgg | attggagggtg | atttgtagat | aatggatagg | 600 |
| tgattttact | cttgcaattt | ggattgaggg | gtggggaaaa | ccagaaaggg | gctggggggg | 660 |
| aaattagtag | aagggtcacct | tgaattcatt | gtggtccata | tcaatgctga | aactgattgg | 720 |
| ggaacttttt | actcttgagt | ccctttgtaa | gggaacccca | gaaag | | 765 |

<210> 216

<211> 780

<212> DNA

<213> Homo sapien

<400> 216

| | | | | | | |
|-------------|------------|-------------|-------------|-------------|-------------|-----|
| cctttttctg | tggcaaatgg | aggtcttttca | ctgcctgtag | agacaatata | gtaagcatag | 60 |
| ttaaggggtg | ggtcagaaca | tgtaagata | acttactgta | tatgtattcc | cttgtatttt | 120 |
| gttaaagctg | gaacatttga | tattttttcca | tttatttatg | aaaaaatatg | aacctatttt | 180 |
| cattttgtaca | aggtaattgt | tttttaaagc | aagtcacctt | aggggtggctt | taattgtata | 240 |
| agtcaagcac | atgtaataaa | ttcaaaacct | gcagttaaca | ggatattaga | catcaatcct | 300 |
| ggtaacccaa | tattaaagat | tctcttttaa | aaagactgaa | catgtttaca | ggtttgaatt | 360 |
| aggctaaaag | gtcttgcagt | ggcttttcat | ggcccttcaa | attggaatgg | aactactgta | 420 |
| ctttgcccatt | tttctataaa | tcagtacttt | ttttttaatt | ttgatataca | ttgtgtgaaa | 480 |
| aaagaaaatg | gctaataaac | tgtattaaat | cttaaaacaat | gtataaagat | tgcaacttagc | 540 |
| cagttcaaag | tgtatactta | ttcataatga | attataacag | ttatatattct | gtgttttctt | 600 |
| gtaaatgttt | cttttccctt | aaatacagat | aattcatttg | tattgcttat | tttattatga | 660 |
| gctacaacaa | aaggacttca | ggaacaagta | atgtattagt | atgggttcaag | attgttgata | 720 |
| ggaactgtct | caaaaggatg | gtggttattt | taaataataa | tagctaattgg | gggttggtaaa | 780 |

<210> 217
 <211> 810
 <212> DNA
 <213> Homo sapien

<400> 217
 ctttttaggca gcccggcacc ttcattccata ggcagagaga gaactgggtg ttggagactt 60
 attcgagggg ataggaaggg ccctgtgaag ttgatttaac ttttggatgt cagactgtga 120
 aagctcctga gaaacttggg gtaataggat cttcttttgg ggatgaaaat ggggaaggcg 180
 tgaggacctg gactacttct ccctagggtca gaaaaagaga attaccctt gacaaatatg 240
 atacctgcta ggtatttccc agggaaattt agggattggc gtctttccct agcatgtgga 300
 ggaattggca gacagcttcc taaggcgagg gagcgggggc ccaaggctga cactgcttgc 360
 atccacgtga ccttaagtta tggcagatga ctctgaaacg gactgaggcc aatgagaaca 420
 gatggatgga gcaactcagg tagacttggt ccttctccta tgctggagga gagggatggg 480
 tctctagaat gttggagggt agttgagagc tcgctctctg aatgttgaac agtgtactct 540
 tctgaaaact gcatattcac tttatgtggg ttcagaatac tgggctcaat actaacataa 600
 gaaagacact tcattgagaa attcttaagc ttacagaaaa cctatctctt tgcacattcc 660
 acataacccc tagcaaatg caggttcttc atacttctgt cctttttcca ttggaagaat 720
 tgcttaagga aaaattaatt cctatttatt cccacaaaag gttgggcatt gctttgattt 780
 taccctatgg gggaatgtgc ctttgaattt 810

<210> 218
 <211> 817
 <212> DNA
 <213> Homo sapien

<400> 218
 ctgctccctt atggagggtc cttcattaat aattattgga tagatagaga aggtgagcct 60
 gtggcttcca agtaccggct tttgctgaag gtctacatgg gaagaagagc atcatttgat 120
 attcagtaga tctgccacac ccaactggct ccatctctg gaaaacagca ctactacaa 180
 gcaactgtaa tagcaccag caatgaccac gctgctctg ctggctcttc cgtacaccag 240
 taaatgaact caccaatgta ttgcacacat acatttcaca gtagtacaat aaagccctgt 300
 atcaggagtg gtaattcaat gacttgactc tatagtgcac tgcagcttta tgtcatacca 360
 acattcaaat attcaaatat ccttccaatc catttgagca aaaatacacc atggctgcca 420
 agacacatgt atttttcttt cttccatgga ctctaaact gctcccacaa tcagcagtgt 480
 tcttctctca gaaattatct taagcttctc tactcaatgg gaggtacaca cagagacctg 540
 agaatatgca gaggccagaa tctctgtctg tgctagagat caactgtact ctgccacct 600
 ggggaacaca tcctctgggt aaagtactcg gaagtaaat acattccctg gagacagata 660
 cgggctttca ctgcagcctg ttagaaaaca caatgtctgt aagttacctc ataggtcaaa 720
 gagttttgga ttatattttt cataatgggg ctatggcctt tttaccctgg ttttaatata 780
 gaaccacctg cagaaaggac attgaaatta aaagcca 817

<210> 219
 <211> 661
 <212> DNA
 <213> Homo sapien

<400> 219
 ggatgctgag gcaggaggat tgagtcctgg agtttcagga tacagtgagc tatgatcatg 60
 ccattgcact ccagcctggg caacagagca agattctgtc tctaagaaaa ggaaaaagaa 120
 aatgaataga tagtggtatt agatgttaat gacatcagtt gtttttattc tttattcttt 180
 cttagaaaca gattagtttt ctcgaattaa agaactacca tttttctttt ttctacaact 240
 ttcaagagct ggtgaagaaa tgatgttttag atttaataga tatagtagca gtcatatatt 300

```
<210> 220
<211> 792
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(792)
<223> n = A,T,C or G
```

```
<210> 221
<211> 759
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(759)
<223> n = A,T,C or G
```

| | | | | | | | |
|-------------|------------|------------|------------|-------------|-------------|--|-----|
| <400> | 221 | | | | | | |
| ctttttctgct | gctccgggag | gtggagtggc | ctggcagagg | gcacatggct | gccacctgct | | 60 |
| gcaaggaaaa | ttctcagtga | agactcctca | gtatgaagga | gataagcctg | cacaatcagt | | 120 |
| cactgataga | tgcttagtgg | aaaaacttcc | aattcccatt | tacagctctc | agagctagga | | 180 |
| ttaaaaaactc | ctgggcataa | actcatgtga | tgagaagtta | tagcacgccc | tcattttcta | | 240 |
| catanccact | tgcatttatg | gttggtttt | gaacttgcta | gaagggaaaag | aagtgc aaat | | 300 |
| gtgtcctcct | tagagctact | ctcctcccct | tggtgggttt | ccagtttgtg | cattgtccag | | 360 |
| atggcccagg | agctgacgat | caaagggaag | aagtcatgtt | tgtcatgaga | atgctttgct | | 420 |
| gcatcaggat | tcagtgaagc | tgttcaccgc | ctggagccca | tgcagcctca | agaggcagga | | 480 |
| tggagctcag | aaaccatcac | tgaqqttaga | aagtgagcac | caaagttgag | ggaagcccac | | 540 |

```

aggagtgagc cgaagtgctc cctttggatt tccaaagtgg gtgctgctgc ttcttccatc 600
agccttgctt ctgaccccaa tgcgttcctg gtgccttctt cttggcattt tgctgtcggg 660
ggcccaagga aaaaaattcc tgcattggcag tggtgaaaaa agatggctgc ctgctgaaac 720
ctgatttggc ctgggtaagc cttttggagc cccgggtaa 759

```

```

<210> 222
<211> 699
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (699)
<223> n = A,T,C or G

```

```

<400> 222
ccttntnaag agttggcatt aattcttcac taaatgtagg agtagaattt atcaggtaag 60
ccacactgac ctctggnctt nttnncgccc gatgattttt aattagttga atccctttac 120
ttgttatata tgtattcata tattctgttc cttcttggat ttacttttat gattggtgcc 180
tattgaggta tttatttcta gtttgtggta cttcatgtgt ttaggttttc tagacagtgg 240
acatagaaga ttcaagaagc taaatgtagg agaatgtnta atgtaggana ntgaggcnac 300
natatcatca atgaatgact tgaagtttcc tctgttgtaa agaatgatat taccataact 360
gccatagnta atattgatgg tgtaagtcaa ataanaaggc aggaggaaag ggacatccat 420
cactgaacca canatcagag nctcattgaa gcctttgaga agaatccaca aaattttaca 480
ggataattca tttctgcgga tcaccacnag aagagaaact ggttaaacag acagggtattc 540
cagagtccaa aaatttacat ttggtttcng aaccaaagac ctcagctccc aggccacagc 600
aaaagggggc ttatgaattc cctggcacc cagnccaaga cccaanaacc tcactctgat 660
tggtttnggg cttgggaaac caaaaaacca atgggtgggc 699

```

```

<210> 223
<211> 598
<212> DNA
<213> Homo sapien

```

```

<400> 223
aaaaagagaa agtttcagat ttgccattca aggccttattt atatatatgt gtgtgtatat 60
aaatacatgc acacacttgc atacatatat atttttggct gggggagtgat gagttttgcc 120
tttctaaggg agggaccgcg caggctcctt tgttctgtat tctggcggag atgggtcctg 180
gccttggtgc actggcctat ccttaaagat catctcccat cctcccagc gccatctgtg 240
tgcagcaacc agaaagggat gaacttgccc ctcttgcggg cctggacaag gtctcttct 300
taccctttct gttgccagtc agcaacctgt aactcacatt ctcttcccag tgaatccctg 360
ggagcgcttg accctgggtg gctgttcagc ttcctgctgc tggggccagc aatttttgag 420
gatttatctt taggccaggc ttgcctcctg acttatccct gctctcccat ttctctcttg 480
tttgagagag aatgaggaag caaagagtga gaaagaatag gggctgaaga cgccactccc 540
agatggctct ttctatctg ctcttctgtt gaaacacacg tgctgtgggc ctcaggcg 598

```

```

<210> 224
<211> 501
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (501)

```

<223> n = A,T,C or G

<400> 224

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| aaacctttat | gatgaattcc | ttatgaatta | ctgaacgaac | actggaatgg | gactcaggta | 60 |
| tcttgaggac | atctctcaac | tctggcctta | gttccccctc | tgtaaaatta | gggtgccaac | 120 |
| taaatgatct | acaaggtccc | ttccagcgcc | gccatttctgt | aattacatca | tgtgtaactg | 180 |
| tattaaacat | acacaagtga | ctgccaggca | tgggaatgta | acttccgagt | aaatgctttg | 240 |
| gtttgttcag | aatacactat | gaacttcttt | ccaaagacgg | gttgtggtaa | atagtggata | 300 |
| ttttgattat | aagaaataga | gtttccttga | agcttttagct | ggagatacag | caatagtgtg | 360 |
| gtgttcctac | aaatatcaca | gtgtattcaa | acatatTTTT | ctatcaaaaa | tcattttttgt | 420 |
| aaaagctgtg | tgtttttatc | caacttgtga | taataaatgt | tctttatttt | agaacaaana | 480 |
| aaaaaaaaaa | aaaaaaaaaa | a | | | | 501 |

<210> 225

<211> 295

<212> DNA

<213> Homo sapien

<400> 225

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| cctgtatagg | gctcgtttcc | ccacacatgc | ctattttctga | agaggcttct | gtcttatttg | 60 |
| aaggccagcc | cacacccagc | tactttaaca | ccagggtttat | ggaaaatgtc | aggaaaaaaaa | 120 |
| aaaaaaaaaa | cacatgcact | cacacaatac | ccaaacatca | raattagaag | ggcataaaac | 180 |
| agggggcctt | ataggctgaa | aaatatctta | ratttcaraa | cagaatacca | atcaaatatt | 240 |
| gaaaattcct | ttgttcaaaa | cacaaagatg | ttttgttttt | aatggggagt | ttttt | 295 |

<210> 226

<211> 372

<212> DNA

<213> Homo sapien

<400> 226

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| agattcctgg | cttagagcat | gcgagcattg | aaggaccaat | agcaaactta | tcagtacttg | 60 |
| gaacagaaga | acttcggcaa | cgagaacact | atctcaagca | gaagagagat | aagttgatgt | 120 |
| ccatgagaaa | ggatatgagg | actaaacaga | tacaaaatat | ggagcagaaa | ggaaaaccca | 180 |
| ctggggaggt | agaggaaatg | acagagaaac | cagaaatgac | agcagaggag | aagcaaacat | 240 |
| tactaaagag | gagattgctt | gcagagaaac | tcaaagaaga | agttattaat | aagtaataat | 300 |
| taagaacaat | ttaacaaaat | ggaagttcaa | attgtcttaa | aaataaatta | tttagtcctg | 360 |
| atgaaatgaa | at | | | | | 372 |

<210> 227

<211> 599

<212> DNA

<213> Homo sapien

<400> 227

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| ggcccccgtc | gcgggagccg | cttcgggcct | tctgggcatg | tctgcatat | ggctccaggt | 60 |
| ttgtttttct | ccccggcact | ctgacgggga | gggtctcccg | catctcctgg | catccgggta | 120 |
| gaggacgcgg | aggatgctga | gctgctggcg | cactgcagca | caactagaga | tgtacggatg | 180 |
| cccccatctt | gatcttacag | aatcagaggt | acagccgcga | gaaagagtca | agaacagaca | 240 |
| gagtcgcttg | aggactcagg | agggtgtttg | ctgcgttgac | aacagactac | accctcacag | 300 |
| tttgctctgc | tcttccaaca | ccagtgggaag | atgatcacat | cccagggatc | agtgtcgttt | 360 |
| agggatgtga | ctgtgggctt | cactcaagag | gagtggcagc | atctggaccc | tgctcagagg | 420 |
| accctgtaca | gggatgtgat | gctggagaac | tacagccacc | ttgtctcagt | agggtattgc | 480 |
| attcctaaac | cagaagtgat | tctcaagttg | gagaaaggcg | aggagccatg | gatattagag | 540 |

gaaaaatttc caagccagag tcattctggaa ttaattaata ccagtagaaa ctattcaat 599

<210> 228
<211> 343
<212> DNA
<213> Homo sapien

<400> 228
aaagtaaatt gtatgaaaaa ttcattttctt caattgcatt agccacattt tgagtattca 60
tgtggctggt agattctgta ttagcacaaa gatatggaac atttccatca ccacagaaag 120
ttctgttgga cagcactgca ttagaatatt ttcatactgc tcttctctca ttaatttttg 180
ttgttaattg tgatgtcttc attggatggg tcataatgtt ccatgaaacc gctcaagtac 240
acaattgtat gttcttttga tcccttacca caaatatctc gctctgctca tttcttttgc 300
agcttcttat aaagtttgtc ttctcaaaa aaaaaaaaaa aaa 343

<210> 229
<211> 417
<212> DNA
<213> Homo sapien

<400> 229
ctcaagctgc agtccaccgg gtatggttct ggatggttcc cccaagggag caggtatgta 60
ggaggtgaag aaaactgaga tttcaagtat gggagagttt ttactatctc cattcctgga 120
ttaaaagtgc tgaaaaagtc cacagttaaa cattccttta ttcacctat ggctcccaag 180
aaaagcattc ttcctctgga gtactgggtg actaagggga caatacacca aatttggtga 240
gtttacaatc aagtctacta aggttggtgact tcttatcag tttggcagag tcccagggca 300
gaataatcat ccactacag gtctctgttt cctctccctc cgcagcagtg gagagcatcc 360
cagtgtttgg ggcactgtgt tctcttctgt ccttgcacca gacctggaa gccttgg 417

<210> 230
<211> 462
<212> DNA
<213> Homo sapien

<400> 230
gaaataccag aagagaaagt ttcattgtgc aaatctaact tcatggcctc gctggctgta 60
ttccttatat gatgctgaga ccttaatgga cagaatcaag aaacagctac gtgaatggga 120
cgaaaatcta aaagatgatt ctcttccttc aaatccaata gatttttctt acagagtagc 180
tgcttgctct cctattgatg atgtattgag aattcagctc cttaaaattg gcagtgtctat 240
ccagcgactt cgctgtgaat tagacattat gaataaatgt acttcccttt gctgtaaaaa 300
atgtcaagaa acagaaataa caaccacaaa tgaaatattc agtttatcct tatgtggggc 360
gatggcagct tatgtgaatc ctcatggata tgtgcatgag acacttactg tgtataaggc 420
ttgcaacttg aatctgatag gccggtctc tacagaacac ag 462

<210> 231
<211> 328
<212> DNA
<213> Homo sapien

<400> 231
ctgtgggttt tctaaacgc ccctcatctg gttgaagccc tagtgtttct ttctcacatc 60
agaggcaaag gcattggggg gggctctggtt tggacaataa atttccctctg gtttggacca 120
agaaaaacag agttctttga ccgctaact atattgtaaaa agaaagtttg taaaaacaag 180
agttaaaatg cttctaacag tgtggtcatc actgcacagg acactggaat tggcattcgg 240

ggttgtgtct gtccatgtgg tttcgttgta tgtcatgtgc tctcagctca gacagagaca 300
tccaattgac ttctgacttg gggcattt 328

<210> 232
<211> 595
<212> DNA
<213> Homo sapien

<400> 232
cgccaatttt agcaaataag agattgtaaa agaagcagat tgaatgaaga attttttagct 60
gtgcagatag gtgatgttgg gatggaaaat gctaatacaac taccctttct tttatcaagt 120
aattaaaata aatctacata aagaaccaa aaggctgttt tataaaagt aaatatccag 180
tatttcagag ggccaggcaa gagcacttca gatgaggcag tcaaaatcat tttttccag 240
tgaggataga ccacaagtgg gtggtgagac cattgaaagc ctttatcaac tgaagagtcc 300
atttaacagc ataatttgtg ggaagactgg aatagggctg aataaatgtg tttgaatctc 360
taattttata ctttcttttc ctgaggaact tgatttttct gtccctggat cgccttgtca 420
taattgggtc tgttctttt actaccactc ttgagtccat atatgaaatc attaaagttg 480
gatgatcagt tttttataaa aatatatatt tttgtccaag aaaaaaaaaa gcatacatat 540
gtgattatgg ctaaatacaa ggtaactgga atgtatatac ttttgctaatt gttcc 595

<210> 233
<211> 600
<212> DNA
<213> Homo sapien

<400> 233
atgaaggtaa actctaaaat cttcataggt caacaaagaa aattttatcct tcacacttat 60
ttctagaaag cagcagggct tatttcttag attgcttaca atgaagctag aatatctgcg 120
ataactgtag agtttcaaaa aggatcccta gggctacttc tacgttctcc ttaccagttg 180
agcactctcc ataatttcca gacgggtcat gggggagaat gatagaaatg agcgtgggaa 240
gaaagacaat gaaattagaa atgggtgaga cacatgggtg tagaatgcta agagcaggggaa 300
tcaggacaat caaccagtg tctaggaagg gtcaagtcac cagtgtcatc tgctgaccaa 360
tgttaggaag aaataaactc aaaggaaaca ccacattttt ccaattaaac tcaaacttat 420
tgacttgtgg tggttctttg atgttgtggg gactgctata acagaaacca attggatttt 480
caagggcaag aaactttgcc actgaataag atgatgtcat ccttctgat aacaaatagg 540
aatgggtggc cagctctaaa cagcgtggac tgagggagtt gcttttctac aatattactt 600

<210> 234
<211> 500
<212> DNA
<213> Homo sapien

<400> 234
aaatttcctaa ttcttttact atcttctcaa cttttcccaa agataaaata aatttcacat 60
aatttcattg aggggaaatg gtagttgtaa aaaactacct caagtagcaa tcaccgttgg 120
cagtgttttc tcactttctg ttctgcaatt gcaatcacac ttccaaaaag aaaagcaaat 180
gtttgctaaa ccatagacag acaacctctt tgtgactggt attataaggt ttataatgaa 240
aacttatcaa atataaaagg tgctccctct tgaaaatgtg tattttatatt gaagttttga 300
gtaagaggtg agtgtttggc aattttcaac actcccctca aaaatctccc aaagttgcaa 360
aaaagtcagt ttagtaaaat tccaagcact taaatgcttc attgagggcc agttgatata 420
cgcaatgcac taatgtgtaa aaattaaccg aatgcaacta ttttataatg gagagctctt 480
accttttct tccagttttt 500

<210> 235

<211> 159
 <212> DNA
 <213> Homo sapien

<400> 235
 aaaattttaca gataaaggca gttcaatact gccactgaga agtacatctc ttaacatata 60
 caacttttcag gccacagttt tgaaggtctg aagtattaag ttggtttgat gaattagtcg 120
 gttggcactt acgaacacat ttattgcctt gccatcttt 159

<210> 236
 <211> 254
 <212> DNA
 <213> Homo sapien

<400> 236
 aaataagtga ataagcgata tttattatct gcaaggtttt tttgtgtgtg tttttgtttt 60
 tatttttcaat atgcaagtta ggcttaattt ttttatctaa tgatcatcat gaaatgaata 120
 agagggctta agaatttgkc cattttgcatt cggaaaagaa tgaccagcaa aaggtttact 180
 aatacctctc cctttgggga tttaatgtct ggtgctgccg cctgagtytc aagaattaaa 240
 gctgcaagag gact 254

<210> 237
 <211> 591
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(591)
 <223> n = A,T,C or G

<400> 237
 tttttttttt tttttttttt tttttttcta atttttactt tttctcaagt ttaatgtara 60
 catacaaraa aacatcaagc aatgtttatt gkgcaattcc aatcattatt tgcaraatct 120
 tggtttaaag tcagtyttta tagccatttc aactgcttgg tttaaacaaa aagcaacaat 180
 ctggttatyt acctataaat ttcatgggat ttytttaaac actgaagtac taaaagcact 240
 gatgatttgt attataattt ttaaaaatatt taaaacctac acagatttca taratcattc 300
 cttttataaa ataatcaaaa taatttgatt atytggaata aaaaattctt gaaacaragc 360
 cctttccagg tatyttcaat ctctgtaaaa ccccaaacc ccaacagagt aratgatgaa 420
 ataaggattt ctcagttgcc caagactgtc tgaaatttaa ggttgaaaaa tggactggcg 480
 tttttcatgt ttcctgngaa ttcanagctt acagggtggc tcaaaactca aatctctggg 540
 atggctttac atggctttca ctttgatttg tttcattttc atttgcttct t 591

<210> 238
 <211> 252
 <212> DNA
 <213> Homo sapien

<400> 238
 aaatggcttt tgccacatac atagatcttc atgatgtgtg agtgtaattc catgtggata 60
 tcagttacca aacattacaa aaaattttat ggccaaaat gaccaacgaa attgttataa 120
 tagaatttat ccaattttga tctttttata ttcttctacc acacctggaa acagaccaat 180
 agacattttg gggttttata ataggaattt gtataaagca ttactctttt tcaataaatt 240
 gttttttaat tt 252


```

gctccacatt tgtagcgaac actttgactc caaagagaag gaggaagaca aagacaagaa      60
ggaaaagaaa gacaaggaca agaaggaagc ccctgctgac atgggagcac atcagggagt      120
ggctgttctg gggattgccc ttattgctat gggggaggag attggtgcag agatggcatt      180
acgaaccttt gg                                     192

```

```

<210> 244
<211> 616
<212> DNA
<213> Homo sapien

```

```

<400> 244
aattttatag caatatactg accatttctaa aaataacaaa atacatgttg ctctcaacta      60
catagttaaa aaaggtagta aattctctta cccaaaatag aggaggggtg ggctagttag      120
ctgctcaaac atttgtaaca aataaaaaatg tatctatata catataatga tcatgttttc      180
atagcctaaa atcaccatac aaaatctaat aataaaattg tgtcgtgttc aggagttggg      240
aagccaacac attaaattaa caaagtattt ttggtatatg taaataatgg gatagaatct      300
ctcgaatcag gattgtccca gaagttctaa ggcagatgtc aatgacatgc acattgtcca      360
tggttcagtaa ttttcaaaga ctagaataaa ctatgtaaac tattcaatac aattcaatat      420
tacttaactg ctaaaaagta cttcaagatc ttgcaactgcc ttgagttagt ataatacaat      480
tagtaattgg aaaatagctg taatagcagg cactgaagaa ttctgacaaa taccaaataa      540
ctgtttgttt ttaccaaata aactggtaag atgatatcac aaagggtttt aagttatttt      600
gctatacaag gttttt                                     616

```

```

<210> 245
<211> 165
<212> DNA
<213> Homo sapien

```

```

<400> 245
ttggaacagt ggattaaaat ccagaagggg aggggtcatg aagaagaaac caggggagta      60
atttcttacc aaacattacc aagaaatatg ccaagtcaca gagcccagat tatggccgcg      120
taccctgaag gttatagaac actcccaaga aacagcaaga caagg                                     165

```

```

<210> 246
<211> 229
<212> DNA
<213> Homo sapien

```

```

<400> 246
tgtactggat ccctccaggt gggggcgact ctcacctgac tattacaata gcctcctaag      60
tggtttccct acttgcaacc ttgcccgtat aatatctatc ctccacacag caggcagggc      120
gatcctttta gaatagaagt tagatcatga aaatgctctg ctctgatccc tgcaaaagct      180
cgccacctcc ttacagtcac cgctgaactc gtagcagagg ttcaggagg                                     229

```

```

<210> 247
<211> 338
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (338)
<223> n = A,T,C or G

```

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ggaacccgtg | tgtacttata | ctggatgatg | ccaccagtgc | cctggatgca | aacagccagt | 60 |
| tacaggngga | gcagctcctg | tacgaaagcc | ctgagcggta | ctcccgtca | gtgcttctca | 120 |
| taccacagca | cctcagcctg | gtggagcagg | ctgaccacat | cctctttctg | gaaggaggcg | 180 |
| ctatccggga | ggggggaacc | caccancagc | tcatggagaa | aaaggggtgc | tactgggcca | 240 |
| tggngcagcg | tcttgcagat | gctccagaat | gaaagccttc | tcagacctgc | gcactccatc | 300 |
| tccctccctt | ttcttctctc | tgtggtggag | aaccacag | | | 338 |

<211> 177

<212> DNA

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| tgaaaacaaa | tgaattctca | actcctacgg | ttcattgtaga | gtttagagaa | aatttcacac | 60 |
| attgtcatca | ttgaactgtg | aacctgggaa | gccagatcat | gattaacact | gacatcaagt | 120 |
| ttcaagttgc | agatcaatgc | acccagtgtt | cagatgaggc | aaacttctcc | gtgacaa | 177 |

<211> 263

<212> DNA

<213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| aaagtaatga | ctttattaat | aaatatacat | ccatatgatg | atgtagatac | aaatcatgaa | 60 |
| cactactcca | ttcccataca | cataattgca | cacgagtagc | tcaagttcat | ggacataaaa | 120 |
| acatacacag | tatctattca | gactttttac | agcagaggac | agcgtgctta | ttatcagtta | 180 |
| attggttaatt | attttctcca | aaattacctg | tggaaaaaag | aaattctgaa | aacttaaaaag | 240 |
| aatcaaagtg | atctgattac | ttt | | | | 263 |

<211> 333

<212> DNA

<213> Homo sapien

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| aaaaaaaaca | acagcgtaaa | tattagccca | caagagcagt | cctaaacaat | cacaattaca | 60 |
| ctgtactacc | caagaagact | gtttattgtg | aagcatttac | ctttcaaaaa | atcattacat | 120 |
| ttctattttct | tgggtggagca | gcacattgtg | gagtgtgatt | cttaattctt | cattgagttt | 180 |
| gtcaatagga | cattgatgct | ggataggttg | tcttttgttt | ttatgcctca | gaccatcttg | 240 |
| tgagattggt | tgctatctc | ataatacagt | tttatgcaga | aaggttgaaa | ctatgtaaat | 300 |
| qgtttttatg | gaaattatca | gttacaatat | ttt | | | 333 |

<211> 384

<212> DNA

<213> Homo sapien

| | | | | | | |
|------------|------------|-------------|------------|-------------|------------|-----|
| aaaccatttg | tacaaaactt | ctataaattt | ttctctctct | ttctctctta | tgtacaaaaa | 60 |
| tatcttaata | tatccccgaa | ctgggttagga | tagatacaaa | tagatttttt | ataataaaaa | 120 |
| attcacaaaa | gattggaagc | attctataat | gaaaatggta | gaaaagacag | tgtgagggaa | 180 |
| gccatggggg | ttgggaatcg | ggccctggag | gagaagcaga | gtttcaaagg | gctgagaata | 240 |
| gcatagtttc | actgtaaacc | aatgtctaca | gcttattggg | gtggggggcta | ctgagacgaa | 300 |

agacaccaac tcgtttctag agggctaaga actgcacttt aagaaagggc ggggaggtga 360
agggacccga gcaagaactt tcag 384

<210> 252
<211> 211
<212> DNA
<213> Homo sapien

<400> 252
aaagcagtct gaaaatggga catctgtaga gaaattcatt tccttcttct cctccggatg 60
tggaatggaa gctttgaggg aaggaaaagt aggaaaagag cgggatggga tgggatggga 120
tgggatggga tgggatagga agagaggctg gggaatgggc agagaagggg gtgctgagtg 180
tgctgtgaga tagagcaaga tcacaagaag g 211

<210> 253
<211> 135
<212> DNA
<213> Homo sapien

<400> 253
aaaaattggt tcttgacaag ctgacttggc acttaagtgc acttttttat gaagaaaaag 60
tacaatgaac tgcttttctt caagcaataa ttgtttccaa cttgtctggg aattgtgtgt 120
ctggttaactg gaagg 135

<210> 254
<211> 361
<212> DNA
<213> Homo sapien

<400> 254
cctgtagccc ctgctacacg ggaggctgaa gtgggaggat cacttgaacc aatgaggggtg 60
aggttacagt gagcccagat catgccacta ctctacaggc tgggtgataa gagtgagacc 120
ctgtatcaaaa aaaaagacaa ggaaaaaaaa aactgggccg tttgtttttg cagaatgtct 180
ctcaattttgg acttttttggg caggaatata atacaagtga tacaaatgct tctttaacat 240
tagaacctgt ataaaattac cattacagac cttgctatct tacttatagg taaatcactg 300
tttaccaagg taagtctttt gggaatttcc aaaaatgaag tccatggaca gttaaaaact 360
g 361

<210> 255
<211> 331
<212> DNA
<213> Homo sapien

<400> 255
aaaaaaaaataa ataatccacc aacgtgattg accttggcga gatcatgttt ctagtctata 60
cctcagtttc cccatctgta aagtgaggat aatgtcccac cccatgtaac tgtggtgagg 120
accaactgca aactgtgcc tgcgagtctc cttggaaaaag tgtaaggttc tacacaaatg 180
gaaagtgatc tgatcacact cagtgtcccc agcccagcct ttcagtgtcc tggccctggg 240
gtggggggaca atactctcct caccctcttc actagtcttc atgaatagca aggaggccat 300
aacataattt ggtctaaacc ccttcctttt t 331

<210> 256
<211> 186
<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(186)

<223> n = A,T,C or G

<400> 256

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| cctttgggcc | cttgcacttt | gacctgcaat | ggggccacac | cagccttgct | tgtgtccacc | 60 |
| tggaaggact | gagggagggt | ggcacgaacc | atgcctgggc | tcaggccggg | cccanagcac | 120 |
| ttgaccttgg | acgcatctgt | cacatcatgc | acaggggacct | tgaaaggact | gcctggcact | 180 |
| tgatgg | | | | | | 186 |

<210> 257

<211> 255

<212> DNA

<213> Homo sapien

<400> 257

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| ctgggggtccg | tcaccgacct | ttggggaact | gggctacggg | gaccacaagc | ccaagtcttc | 60 |
| cactgcagcc | caggaggtaa | agactctgga | tggcatcttc | tcagagcagg | tcgccatggg | 120 |
| ctactcacac | tccttgggtga | tagcaagaga | tgaaagttag | actgagaaa | agaagatcaa | 180 |
| gaaactgcca | gaatacaacc | cccgaaccct | ctgatgctcc | cagagactcc | tccgactcca | 240 |
| cacctctcgc | ggcag | | | | | 255 |

<210> 258

<211> 604

<212> DNA

<213> Homo sapien

<400> 258

| | | | | | | |
|------------|------------|-------------|-------------|------------|------------|-----|
| ctgaatttgc | aatggagttt | ggtggtgcaa | tcggtattga | ttagtttggc | atagacagat | 60 |
| gcagcagttt | agagcaaaat | cgagaaaatg | atctctctga | ttctccttga | tttcctggca | 120 |
| gaagatatct | tactttttca | gcaaaactttt | cttttaacac | taaagcagcc | tagggcaatg | 180 |
| ccagatactt | agagcttttc | tcttgattat | aagtagaaat | gggggtgtct | gggctagagg | 240 |
| tggagggtgg | atgtgctgtc | gtcacagtct | agctggcagc | aagcaaggca | aaagcagaga | 300 |
| ctgctctaga | agcgggtcca | agcagcagag | acgtcaggaa | aggcacttct | tagtaccac | 360 |
| ctctatgctt | taatagttgc | ttgttaagct | gcttcatggg | ttgagacaaa | ctaccagcac | 420 |
| ttcaaagagc | tcagttctct | gctcaactct | cttctctagt | tacattattt | tttttccttc | 480 |
| aggagactga | ggcaggaaaa | tcgcttgaac | tcaggagggtc | gaggccgcag | tgagccaaga | 540 |
| tcacaccacc | gcactccagc | ctgggccttg | caaagtgtct | ggattacagg | aatgagccac | 600 |
| cagg | | | | | | 604 |

<210> 259

<211> 429

<212> DNA

<213> Homo sapien

<400> 259

| | | | | | | |
|-------------|------------|------------|-------------|------------|------------|-----|
| aaaaatgtct | gtatcgagat | cttccagttt | gaagtcttcc | tcctctgtgt | cttcccaagg | 60 |
| ctctgtggca | agctccactg | gttctcccg | ttccatcaga | accactgact | tccacaatcc | 120 |
| tggtatatccc | aagtacctgg | gcacccccca | cctggaaactg | tacttgagtg | actcacttag | 180 |
| aaacttgaac | aaagagcggc | aattccactt | cgctgggtatc | aggtcccggc | tcaaccacat | 240 |
| gctggctatg | ctgtcaagga | gaacactctt | tactgaaaac | caccttggcc | ttcattctgg | 300 |

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| caatttcagc | agagttaatt | tgcttgctgt | tagagatgta | gcactttatc | cttccctatca | 360 |
| gtaactgctc | cggtgttcaga | ctcctgggtt | cttccaggct | tacagtggac | atcatcagct | 420 |
| tcttgcttt | | | | | | 429 |

<210> 260
 <211> 385
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(385)
 <223> n = A,T,C or G

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| <400> 260 | | | | | | |
| ctgcaacaca | tgcagcacca | gtctcagcct | tctcctcggc | agcactcccc | tgctgcctct | 60 |
| cagataacat | cccccatccc | tgccatcggg | agcccccagc | cagcctctca | gcagcaccag | 120 |
| tcgcaaatac | agtctcagac | acagactcaa | gtattatcgc | aggtcagtat | tttctgaana | 180 |
| cgcataatggc | agacggatgt | gcgtatacca | aggagagtgg | cataggaggg | aaaagcatat | 240 |
| gtggctgaaa | cctgtaagtt | ggtgttggtt | atgcagaaat | gtgtaacaga | tcaaacgggtc | 300 |
| ctctcaagtg | tctattanat | aggcaataag | aactgcagtg | tagctgagta | acatctttta | 360 |
| gctgactata | aatcactttg | ttttt | | | | 385 |

<210> 261
 <211> 230
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| <400> 261 | | | | | | |
| ctgtactgga | tccctccagg | tgggggcgac | tctcacctga | ctattacaat | agcctcctaa | 60 |
| gtgggtttccc | tacttgcaac | cttgcccgta | taatatctat | cctccacaca | gcaggcaggg | 120 |
| cgatcccttta | agaatagaag | ttagatcatg | aaaatgctct | gctctgatcc | ctgcaaaaagc | 180 |
| tcgccacctc | cttacagtca | ccgctgaact | cgtagcagag | gttcaggagg | | 230 |

<210> 262
 <211> 198
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(198)
 <223> n = A,T,C or G

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| <400> 262 | | | | | | |
| atgttaagta | aacatgaaat | ctatataaca | gaacaaaaat | tcactcttat | gtcaatgtca | 60 |
| gcgtgttaat | gtagatctat | ttactganac | agactctgta | gtggcagaga | gtggccttgt | 120 |
| taagccagga | ccctgttctg | caggctgtgg | gtagaagcta | ggaagtccct | ggagtttcac | 180 |
| ccagcttttc | catgaatg | | | | | 198 |

<210> 263
 <211> 157
 <212> DNA
 <213> Homo sapien

```
<210> 264
<211> 290
<212> DNA
<213> Homo sapien
```

```
<210> 265
<211> 234
<212> DNA
<213> Homo sapien
```

```
<210> 266
<211> 335
<212> DNA
<213> Homo sapien
```

```
<210> 267
<211> 619
<212> DNA
<213> Homo sapien
```

<400> 267
tggagctctc acgaagggat cggggaggtg ctggagaagg aagactgcat gcaggccctg 60

| | | | | | | |
|-------------|------------|------------|-------------|-------------|------------|-----|
| agcggccana | ttttcatggg | catggngtcc | tcccagttacc | aggccccggct | ggacatcgng | 120 |
| cgcttcattg | atgggcttgt | caacgcctgc | atccgctttg | tctacttctc | tttggaggat | 180 |
| gagctcaaaa | gcaaggtgtt | tgcanaaaaa | atgggcctgg | agacaggctg | gaactgccac | 240 |
| atctccctca | cacccaatgg | tgacatgcct | ggctccgaga | tccccccctc | cagccccagc | 300 |
| caagcaggct | ccctgcatga | tgacctgaat | caggtgtccc | gagatgatgc | anaagggctc | 360 |
| ctctcatgg | aggaggagg | ccactcggac | ctcatcagct | tccagcctac | ggacagcgac | 420 |
| atccccagct | tcctggagga | ctccaaccgg | gccaagctgc | cccgggggat | ccaccaagtg | 480 |
| cgggccccacc | tgcagaacat | tgacaacgtg | cccctgctag | tgcccccttt | caccgactgc | 540 |
| accccanaga | ccatgtgtga | gatgataaag | atcatgcaan | agtacgggga | ggtgacctgc | 600 |
| tgcttgggca | ncctctgcc | | | | | 619 |

<210> 268

<211> 147

<212> DNA

<213> Homo sapien

<400> 268

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| cctataaccc | agacaccagc | atggacaaaa | ctcagttata | ctgaattcag | agacaaaatt | 60 |
| cagtgcact | cttctaccac | ttatttagg | ttctacagca | tttctactgag | cagacttagt | 120 |
| tttttgtttt | tgttttacaa | acctttt | | | | 147 |

<210> 269

<211> 325

<212> DNA

<213> Homo sapien

<400> 269

| | | | | | | |
|------------|------------|-------------|-------------|-------------|------------|-----|
| ctgagctgta | ggaatgggtt | cttgggtacac | aagatagttat | tggttgagcta | gttttcgagc | 60 |
| tctgtgcaca | agcactctgt | aatcggggcc | catgccactg | tacaccaaac | ctatatgctt | 120 |
| ggtaattgg | tctactttgt | gtacacttcg | ctcatcatac | agaatggatt | tctgtttttt | 180 |
| ctcagttgct | aataccacac | catttgcagc | tttaattccc | acggacgggg | ctcctccagc | 240 |
| tacagcagcc | aaagcatatt | caatctggac | aagttttacca | gacgggctga | atgtagtcag | 300 |
| cgaaaagctg | tacccgcgct | ccgcc | | | | 325 |

<210> 270

<211> 428

<212> DNA

<213> Homo sapien

<400> 270

| | | | | | | |
|-------------|------------|-------------|------------|------------|------------|-----|
| aaacatatgg | taaattaccg | agtgacacct | ctgggctaga | gacctctttt | gaggggagtt | 60 |
| tgcaaaactac | ggattcaatt | tctttaacag | ttatgaagtt | ctttaaagaa | cctgtttgg | 120 |
| attgggggg | tgtggtcacc | tgtgcttttc | tgagattttg | cccctacatc | taagttgttg | 180 |
| aatgcatgtg | tgtagagttg | tttatggtgc | ttccctttct | tcttagaagg | gtctatagta | 240 |
| atatccccctg | ccttatccct | agtagtacta | atttgtgttt | tcttacttct | tgacaggcaa | 300 |
| acacatcaga | gcataagtgg | ttcctaattgc | caagctgacc | tcccttgatc | tctgtcttct | 360 |
| acaggatatt | gacatgggac | ttctttatta | ccttttcagt | tactgatac | cttcaaatag | 420 |
| ctttattt | | | | | | 428 |

<210> 271

<211> 206

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(206)
 <223> n = A,T,C or G

<400> 271
 cgtcccggag cccacgngg ncatggctgg canagcgctc tgcattgctgg ggctggctcct 60
 ggcttggctg tccctccagct ctgctgagga gtacgtgggc ctgtctgcaa accagtgngc 120
 cgtgccagcc aaggacaggg tggactgcgg ctacccccat gtcaccccca aggagtgcac 180
 caaccggggc tgetgctttg actcca 206

<210> 272
 <211> 83
 <212> DNA
 <213> Homo sapien

<400> 272
 ctggcttccc tgagaactca acaatgcctt ttcttgaggg ccttcctcga tcatccacaa 60
 tgactacagc cctctctacc tgg 83

<210> 273
 <211> 472
 <212> DNA
 <213> Homo sapien

<400> 273
 ctggagaagg tgtgcagggg aaaccctgct gatgtcaccg aggccagggt gtctttctac 60
 tcgggacact cttccttttg gatgtactgc atgggtgttct tggcgctgta tgtgcaggca 120
 cgactctgtt ggaagtgggc acggctgctg cgaccacag tccagttctt cctgggtggc 180
 ttgcccctct acgtgggcta cccccgctg tctgattaca aacaccactg gagcgatgtc 240
 cttgttggcc tcctgcagga ggcactgggtg gctgccctca ctgtctgcta catctcagac 300
 ttcttcaaag cccgaccccc acagcactgt ctgaaggagg aggagctgga acggaagccc 360
 agcctgtcac tgacgttgac cctgggcgag gctgaccaca accactatgg ataccgcac 420
 tcctctctct gaggcggac cccgccagg caggagctg ctgtgagtc ag 472

<210> 274
 <211> 205
 <212> DNA
 <213> Homo sapien

<400> 274
 ccaggcggcc cgaggactta cggtcggcac ttctctgttc tcccgtgtca gcggtgtggtg 60
 tcgcttgcac gggctgtacc tggatggtgt gtccaccatc gacacggagg ggctggattt 120
 gtttctcagg caatcctgta ttttaatttt agatgtattt cctgaagcat atttttcata 180
 gaatgtagcg tgtaaatagc ttttt 205

<210> 275
 <211> 308
 <212> DNA
 <213> Homo sapien

<400> 275
 ctctctgccc tccccaccga catcatgctc cagttccagc ttggatttac actgggcaac 60
 gtgggttgaa tgtatctggc tcagaactat gatatacaca acctggctaa aaaacttgaa 120

gaaattaaaa aggacttggga tgccaagaag aaacccccta gtgcatgaga ctgcctccag 180
 cactgccttc aggatatact gattctactg ctcttgaggg cctcgtttac tatctgaacc 240
 aaaagctttt gttttcgtct ccagcctcag cacttctctt ctttgctaga ccctgtgttt 300
 tttgcttt 308

<210> 276
 <211> 201
 <212> DNA
 <213> Homo sapien

<400> 276
 aaattaactt tttcttgcaa aatattcatt tcattttttc caagaaaatc ttataaaggc 60
 aaaaataaaa ttttattttg gcaaatgtca tgaagtcgat actggcagca tatggagtta 120
 gttaaaaata gacaacaact gctagatata ttcaaaattc tatttttttt tctgagcata 180
 gtcaaagaga aatttttcatt t 201

<210> 277
 <211> 520
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(520)
 <223> n = A,T,C or G

<400> 277
 aaaaaaaaaag tattcagcac catttgetca tnggtctttc agagtttggt cttaaagttt 60
 ctggaacttt cctgtctgta aagtaacagg aattactgag ctacattgga aagcctctct 120
 gggacaggca gtggggagtt aagcagtcac cataaaggaa tcagtgtaca ttcagcatgg 180
 tgacttgact acacaacaat cccttccctt ctactgtagc tcaagagaga catgcttcta 240
 accactgagg tatgaggagt ctccagactgt tatttgctgt tagaattggg cttcccagct 300
 aataacagta catctctggc acagatgcta ttggtcctta atgtcctgtg attttaggaa 360
 atagtttgga tttagttcaa tttattcaga aaccaaactg gtttaattag cttcactact 420
 ctggcagagt aagggtatgc tggtttagta tctttataaa atatatataa tgttataggta 480
 aatcatagtc ttaaatcata cctaaaatac tgtatcattt 520

<210> 278
 <211> 264
 <212> DNA
 <213> Homo sapien

<400> 278
 cgcgcggggc ggaactttcc agaacgctcg gtgagaggcg gaggagcggg aactaccccg 60
 gctgcgcaca gctcggcgct ccttcccgtt ccttcacaca ccggcctcag cccgcaccgg 120
 cagtagaaga tgggtgaaaga aacaacttac tacgatgttt tgggggtcaa acccaatgct 180
 actcaggaag aattgaaaaa ggcttatagg aaactggcct tgaagtacca tcctgataag 240
 aacccaaatg aaggagagaa gttt 264

<210> 279
 <211> 414
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaacatacaa | taatttttat | tatggaaatt | aatctttaca | tacaaaatca | gctacgtaat | 60 |
| tttacttaca | aaacaataaa | aactgttctt | tactgtggca | acaaaagaag | cattttgaca | 120 |
| aatgaaaaaa | attaatgcaa | acaaattaaa | acaatgcttt | tctttttact | tgtttcactg | 180 |
| tctcttctat | ttattttcta | tgatcatttg | acacaaacat | ggattacttt | gatatctact | 240 |
| gaaacataaa | tgataagggt | cttaaagggt | gaattaaaag | tctgggtggt | caatatttta | 300 |
| gaagctgaat | aaacaaaacg | aaattggggg | ttgtgattac | agaggattta | tcattttttc | 360 |
| cctttgtcca | tatgaaaata | tataatagaa | aattaccac | gggaaaacat | tttt | 414 |

<211> 262

<213> Homo sapien

| | | | | | | |
|-------------|-------------|------------|------------|-------------|------------|-----|
| ccaccatgcc | tggcctgctt | caatTTTTTt | atgccacttt | gtaaacggca | cttaattatg | 60 |
| gaaaatagga | aaaagcaaaa | ctaaaataag | gaagaggata | tatatataac | ttttcacaat | 120 |
| ctctttttctg | atcccccttta | gatgccagtt | caaccaggac | cacacacaga | tttcatttta | 180 |
| tttgtagagt | atatgaaaag | atttaatatg | ctcatgcatt | ttatttttacg | tatactgatt | 240 |
| tctacgtttt | gactgactat | tt | | | | 262 |

<211> 349

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| ctgtgaccgc | ggtgcatcag | tggatatagt | tgtgtctccc | catggggggt | taacagtctc | 60 |
| tgcccaagac | cgttttctga | taatggctgc | agaaatggaa | cagtcattctg | gcacaggccc | 120 |
| agcagaatta | actcagtttt | ggaaagaagt | tcccagaaac | aaagtgatgg | aacatagggt | 180 |
| aagatgccat | actgttgaaa | gcagtaaac | aaacactctt | acgttaaaaag | acaatgcttt | 240 |
| caatatgtca | gataaaacca | gtgaagatat | atgtctacaa | ctcagtcggt | tactagaaaag | 300 |
| caataggaag | cttgaagacc | aagttcagcg | ttgtatctgg | ttccagcag | | 349 |

<211> 381

<212> DNA

<213> Homo sapien

<221> misc feature

<222> (1) ... (381)

<223> n = A, T, C or G

| | | | | | | |
|------------|-------------|------------|-------------|------------|------------|-----|
| aaacactaaa | tgaagcttct | cacaatttct | aattataaac | aaaaggtga | aaacagtatg | 60 |
| ggaaacaaag | tttcaaaaca | aagaaaagtt | gagtaaaagg | tgccccctct | atggtctatc | 120 |
| tgaaagaaac | attttactca | gagaggcaaa | cattttctgat | ctaggagtaa | gtttccact | 180 |
| cactttgcaa | ggaccctactc | attctgcana | aagacctaca | agtctttctg | gtctcaattg | 240 |
| caaagtacgt | gaaaatgtgt | atgaaagatc | taaaagctaa | atattagaat | aaggctaatt | 300 |
| gaaatcaaaa | ttgtgtgctg | gtctaaatat | acatcttcgg | cttcttctct | tttagtaagt | 360 |
| atttttattt | caqatqtatt | t | | | | 381 |

<210> 283

<211> 543
 <212> DNA
 <213> Homo sapien

<400> 283
 aatatagctc ctccctaccc ccaacaatgg accctgcccc ttgcctccca gttccttgat 60
 cttcctagggt tccacaactc tcttttttcc ttttagtttta ttccctccag ccaaacctct 120
 cttattcaat attttgagcc aatgggggag ttatgtagat ttttttccct acacattagc 180
 tggccctttt tatgaccaat gactcataag gcaagatgtg tgggtggcatc ttcggacagg 240
 cagcaggctt taatagggca gcctggggtg gtggaggcaa gcaaagctaa ttggcatgcg 300
 tgggaatcaa accccaggcc ctgggctcat tagcccatgg tcaaaacaac tgagccagag 360
 gaggtataaa tttgcccagg aatatcagta gttcctttat tagaagaaaa tggctgatat 420
 ggaagttagg gaactctgaat tgccagagaa tcttggggaag agtaataagc tcttagtctc 480
 aacaaaaagt gttttttcat ctccagcgct aaagggtgct atatgggaac aaagaagtat 540
 ttt 543

<210> 284
 <211> 147
 <212> DNA
 <213> Homo sapien

<400> 284
 aaactggtat tttatctttg attctccttc agccctcacc cctggttctc atctttcttg 60
 atcaacatct tttcttgctt ctgtccctct ctctcatctc ttagctcccc tccaacctgg 120
 ggggcagtgg tgtggagaag ccacagg 147

<210> 285
 <211> 316
 <212> DNA
 <213> Homo sapien

<400> 285
 cgcccgaggt ctggcttcac tctactccc tctctgctcg cagcacgtcg gccgccagct 60
 ctttgatgtg ttcccaggcc cgctgcacat gggcagattc caccgtgcga gaacagatgg 120
 caaagcgcag gacaaacttg tccctgaggt gacatggaac caagtggatt tttttggcac 180
 tgtttattct ttgcagaaga gcttcattca ctttggttga accctttagc cgaaagcaga 240
 caagccccag aatgacttcc acacagattt caaagcgggg atcctggcgc accagtgact 300
 caaactcatg ggacag 316

<210> 286
 <211> 322
 <212> DNA
 <213> Homo sapien

<400> 286
 cctggggagc ccttttagtg ggtgggacct caggcagacc cccaaaccaa agggagccag 60
 atgcccaggt tcaagtcatt agtgatatgt ggcagggtg acagagaaat aatcctggag 120
 gtctccaaag ctgctgggaa tggaatggcg atgaaaagcg caggagtggg cagggtgtgg 180
 tgggtgatgg tggcctcact cagagtggac caaggcccca gtccttgcc caaaaccaa 240
 gcccttgggc ccgaagtttt tagcataaca tcctttgcag taaatctcgc catccttgtc 300
 tgccagggtg gttgactcaa gg 322

<210> 287
 <211> 364

<212> DNA
<213> Homo sapien

<400> 287
ctgcccacgc tcaaaccaat tctggctgat atcgagtacc tgcaggacca gcacctcctg 60
ctcacagtca agtccatgga tggctatgaa tcctatgggg agtgtgtggt tgcactcaaa 120
tccatgatcg gcagcacggc ccaacagttc ctgaccttcc tatcccaccg tggcgaggag 180
acaggcaata tcagaggctc catgaagggtg cgggtgccc cggagcgcct gggcaccctg 240
gagcggctct acgagtggat cagcattgat aaggatgagg caggagcaaa gagcaaagcc 300
ccctctgtgt cccgagggag ccaggagccc aggtcagggg gccgcaagcc agccttcaca 360
gagg 364

<210> 288
<211> 261
<212> DNA
<213> Homo sapien

<400> 288
aaaattataa ctactcattc tttcttttagc cttagttaat ttgagcagaa gccacaacaa 60
gcaaaccaca ataaatttag aattggcaga aatccacatt aactcctctt cccaagtttc 120
cacactacta ccatttacag ttgtagggtt gtaatgtata attatgtaat gcagaaacta 180
gctttgactt gtgtaacgat gcactgtcaa agtaagcaaa gtaagaattg aaattccaca 240
ttcccagaat ttaacactca g 261

<210> 289
<211> 261
<212> DNA
<213> Homo sapien

<400> 289
ctgagtgtta aattctggga atgtggaatt tcaattotta ctttgottac tttgacagtg 60
catcgttaca caagtcaaag ctagttttctg cattacataa ttatacatta caaacctaca 120
actgtaaatt gtagtagtgt ggaaacttgg gaagaggagt taatgtggat ttctgccaat 180
tctaaattta ttgtggtttg cttgttgtgg cttctgctca aattaactaa ggctaaagaa 240
agaatgagta gttataattt t 261

<210> 290
<211> 92
<212> DNA
<213> Homo sapien

<400> 290
ccactacccg aacttacagg tgccaaaaga agaaagggtg taaacggaga ccacctatca 60
ctcatcagaa ctaggatca tcacattcct tt 92

<210> 291
<211> 287
<212> DNA
<213> Homo sapien

<400> 291
ccatggctcc gctcagggcc ccggtcacct ccgagtcact ctgttccttg actgtctttg 60
tgtttctgta cctcaaggca ctgaagctgg aggactctgt ccatgcctgt gtcaccctcg 120
tgtgggagcc tctgggctcg gcaggtccac atttcatgag ctgaggcgtg ggccagggcc 180


```
<210> 292
<211> 270
<212> DNA
<213> Homo sapien
```

```
<210> 293
<211> 333
<212> DNA
<213> Homo sapien
```

```
<210> 294
<211> 123
<212> DNA
<213> Homo sapien
```

```
<210> 295
<211> 311
<212> DNA
<213> Homo sapien
```

| | |
|-------|-----|
| <210> | 296 |
| <211> | 241 |
| <212> | DNA |

<400> 296

<210> 297

<211> 295

<212> DNA

<213> Homo sapien

<400> 297

| | | | | | | |
|------------|-------------|------------|------------|-------------|------------|-----|
| aaacacaaga | tgaaaaatact | ctgttctgtc | caaagcatca | cctaattggtg | tgaggcatct | 60 |
| cacttagctg | tggagaagtc | cttggaatta | gatctcagaa | agacagcttt | aagacagtaa | 120 |
| aacctttttg | caatgggcta | attgccttaa | aagaagagtt | ctacctgaaa | gaccttgcag | 180 |
| gtggagaaat | tgtcctacaa | agattcttgg | atatgttagt | ggagataact | gacatgggta | 240 |
| gctgtgggtc | aaccaggaac | tgtcaacaac | ctgatctctg | caaaaccagg | atgga | 295 |

<210> 298

<211> 347

<212> DNA

<213> Homo sapien

<400> 298

| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|----|---|---|-----|---|---|---|---|-----|-----|
| c | a | a | a | a | t | a | a | | g | t | t | c | a | g | g | c | a | a | a | g | | a | t | c | c | a | g | t | g | g | a | | a | t | a | t | g | g | g | g | a | g | | 60 | | | | | | | | | |
| a | c | c | a | a | c | a | c | c | t | | g | c | t | a | c | c | c | c | a | g | | a | g | a | g | c | t | t | t | c | | t | a | a | a | a | a | a | a | g | c | | a | a | g | a | a | g | c | a | g | | 120 |
| t | a | t | t | c | a | c | c | c | t | | g | c | a | g | a | a | g | a | c | a | | c | g | g | a | a | g | g | t | a | c | | t | g | a | g | t | t | t | g | a | | c | c | a | g | a | g | g | a | c | | 180 |
| t | g | t | a | a | a | g | a | a | | g | g | g | t | t | t | g | c | t | | a | c | a | t | c | c | c | g | a | c | | a | g | g | a | a | g | a | c | t | | a | g | c | c | c | a | t | a | t | a | | 240 | |
| a | a | c | a | a | c | c | a | t | g | | g | c | a | a | c | t | c | g | g | a | | c | c | a | g | c | c | c | c | c | | c | c | t | g | g | c | t | g | c | a | | c | a | g | a | g | t | t | a | | 300 | |
| a | c | t | g | a | g | t | c | t | c | | g | g | c | a | a | a | g | a | a | | a | t | c | t | t | g | c | a | | g | t | c | t | c | c | a | a | | c | c | a | a | c | a | g | | 347 | | | | | | |

<210> 299

<211> 268

<212> DNA

<213> Homo sapien

<400> 299

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| aaaaagtaaa | catgaaaaca | tcacgaattg | taccatgatt | caagaataac | ttttgtaata | 60 |
| gaaaacacat | gaccttttgc | agtatagtgt | gataccgaag | taaaagtgaa | agaaataaat | 120 |
| gcaggaaaagt | ttaagtggat | gtaagttttt | ataaggaaag | taataagagg | aggctgcttt | 180 |
| tgaaggtcct | ttgatcttcc | atgatgataa | tatcgttgca | aagttcttta | acttgatttc | 240 |
| aagtaattag | cagttgacca | cttggttt | | | | 268 |

<210> 300

<211> 185

<212> DNA

<213> Homo sapien

<400> 300

| | | | | | | |
|------------|-------------|--------------|-------------|------------|------------|-----|
| aaattggaga | aggaagtttt | cctgaagagc | cagaatcctt | gctaagtcac | ttagatccaa | 60 |
| ctgaccatct | ttattttctgt | caaaaaatcctt | catcatgggtg | cgggtgtatt | cttccagttt | 120 |

agcctcagaa atggcctttc tgtggtgaag aaagaggctc cggaggaagt tgcggagctc 180
agcag 185

<210> 301
<211> 75
<212> DNA
<213> Homo sapien

<400> 301
aaaattggaa agtgggataa gaaatctaaa gtaaccagct tatctttgaa acaatattat 60
tttgaaattg gcttt 75

<210> 302
<211> 247
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(247)
<223> n = A,T,C or G

<400> 302
ccatgtttctc tgaattgggt gcagaagaca agggcagagt ggctgcggcc cctattacct 60
ttgtagcagc cacatcagaa agcagaagaa aacagtatct ctgaaggcat tgtttgaggt 120
tgatctcagc actgaacgat ttcaagccct acgcaccana acagaaggag ggtggaggaa 180
gtgatcanag ggaacgagct gtaggtttgc anaaatgtgt gaaaccaaaa tgatcactgc 240
ctacttg 247

<210> 303
<211> 535
<212> DNA
<213> Homo sapien

<400> 303
ctgcttcaga ggaaatcact gaaaaataaa gaaaaacat ccatgcatgg ctgcatccag 60
tgtacctgta atcctgaaga aaaggctccta attccttcca tgctgaaatg ctagctttgg 120
tttcagagag agactttatt gcaactgtga ccaccgtcac tggtagcac tgctgttcgg 180
ccccagcgg acttaaaaga ctggaatgtg gtagtgccgg tcgttctcgg tcagcaggga 240
gatctccggc cagtccttga gaggctcctc tgggtagcag acttcaaagt ctctggagtt 300
aaacttgaac agtctgaaca cttttatctt tacttcaagg gagtatccaa gtataaacat 360
atcaatctgc tctagtccac atgtgtcggc tacagaattc aggtgattca tcatgaagct 420
caaaggatca gaggatgtct ccctggaaaa caggagtcta aaaagactgg gaatgacctt 480
tttagtcttc atttgttcat aaacttcagt gacttgatac agcatgatga acttt 535

<210> 304
<211> 522
<212> DNA
<213> Homo sapien

<400> 304
ccgcgctcgg tctacaatca cgttttatta ttggctcgtc tagtcatggg atagagaagg 60
taaatagcaa aatagaaaga aaagggggaa aaggtagaag gcaaggggaa aactattggg 120
tttagatctt tatcctgggtc ctgtcaatga tcaggtaatt ggaaggatca aaattaggcc 180

```
<210> 305
<211> 165
<212> DNA
<213> Homo sapien
```

```
<210> 306
<211> 294
<212> DNA
<213> Homo sapien
```

```
<210> 307
<211> 181
<212> DNA
<213> Homo sapien
```

```
<210> 308
<211> 179
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|-------------|------------|------------|-------------|------------|------------|--|-----|
| <400> | 308 | | | | | | |
| aaggtctgagg | actgctggga | gctcagatca | gcccggagct | actggctcat | gggcagccaa | | 60 |
| aaaatactgg | atctgctgaa | cgaaggctca | gcccagagatc | tccgcagtct | tcagcgcatt | | 120 |
| ggccccgaaga | aggcccanct | aatcgtgggc | tggcgggagc | tccacggccc | cttcagcca | | 179 |

<210> 309
 <211> 129
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(129)
 <223> n = A,T,C or G

<400> 309
 ctgcccgttt gcccgtagct gactcagntt cctcatcttc atctccatcc tcttctcac 60
 catcaccttc ttcttctctc tctcttctct cccaccttc ttctcttctc tcgtctacct 120
 cattgtcag 129

<210> 310
 <211> 390
 <212> DNA
 <213> Homo sapien

<400> 310
 tgaggctggg ggagagccgt ggtccctgag gatgggtcag agctaaactc cttcctggcc 60
 tgagagtcag ctctctgccc tgtgtacttc cggggccagg gctgccccta atctctgtag 120
 gaaccgtggg atgtctgcat gttgcccctt tctcttttcc cctttctctgt cccaccatac 180
 gagcacctcc agcctgaaca gaagctctta ctctttctta tttcagtgtt acctgtgtgc 240
 ttggtctgtt tgactttacg cccatctcag gacacttccg tagactgttt aggttcccct 300
 gtcaaataac agttaccac tcgggtcccag ttttgttgcc ccagaaaggg atgttattat 360
 ccttgggggc tcccagggca aggggttaagg 390

<210> 311
 <211> 355
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(355)
 <223> n = A,T,C or G

<400> 311
 cctctctgtg ctgctgaagg cagatcgctt gttccacacc agctaccact cccaggcagt 60
 gcataatccg ctgttgagaa atgccgtgtc tagattgtgg acaagagcct gogtgattat 120
 gctatangga naaaaattct tcgagtcca ccnancctc tctaaacatt tggctcactc 180
 aaaacaaaaa gncaccaatc ttantactgc tgaacttcat ttatgttnacc taacattaac 240
 cntcgtagga aaaccaaata gccctctcgt ncangatatg ttgctaaagg actacnttgt 300
 tcaacacaac ggctccggtg tgtgaactcc tgtttgggtg attcccctac tctca 355

<210> 312
 <211> 498
 <212> DNA
 <213> Homo sapien

<400> 312

```

ccattctttt gaatctaata tattatcaat agcatcctcc ataatatctt tgataaaaagg      60
tgtccaccga gagagctgaa aagtttcttc tgcagaccga tcctttctta acggtttgcc      120
ttgttgagat tggggaacaa tgggaacacc aaggtaactc cagttacgaa tcatgtcact      180
ctcattttct atctttacat tctggatcaa cctgtccaaa ttttcttccg tagttccatt      240
aatactgaag atataaagta gaattgctct tattttatca caattatcat gatttttgtt      300
gagtagaact ggaaggagta ctgcacatgga atctttcacc ttctgtcctt ctgcatcagt      360
tccaagtgcc aggtcctgtt cagttttgca gagcttttct atattaagct tgaacttatt      420
catgcaatct tctgctaagt taagatggac aacttgctta gtaatctgtt ttcggaaata      480
gggcatcttt ttcacatcag

```

<210> 313

<211> 653

<212> DNA

<213> Homo sapien

<400> 313

```

aaacttatca gattttttta agttaggtaa tttcaatcca cagtggctcc atatgggttaa      60
aaaaacaaaa acaaaaacgc atttaaggat acacgaagca gtgaaaacaa agccccagta      120
ttttcgctaa agtactggaa atacctgttt ctaaaaacag ctttatattt gtccactgcc      180
tagaatagct ctcacccaaa cctcaaaaat aagagcagat agattttaga agcaagaaaa      240
ggtaaacagt gcccatatta tttgagactg gctctgctgc cctccctaag ccagtttaca      300
ttctttgaga ttcttgaggt ggggtgagtc ggggtgaaga ctgcacaggc catgtcccct      360
gctccaacta ttctcagaa cgtcccagggt ggagggagtg gcctgtcgat tttcactcat      420
tccatggagc tctgtgtaca tgaaaattcc tccaagtgtg gcttttgtcg aattcagaga      480
tacagcaagc cacgcataaa acatggagtg tagagcactg gtgtacctag cttagaaaca      540
ccctcgggtga atgtggtact gtggctcgaa aggaagcaag ggacaggacc caggagactg      600
ggcgccaggt ctctcggagt tccacacaca cctgtgaagc ccggccagca cag

```

<210> 314

<211> 513

<212> DNA

<213> Homo sapien

<400> 314

```

ctggaagatt ttgctgcatt tggcattata ctgtaattta cagtatacaa catctgggga      60
ctcagtacta tcttagcaca gactaacttc tcccactccg tcagaggtgg cagggtggcg      120
gtcgggtggg agggcctttt ctcccataa atgcctgaac tttaatttat accatataag      180
aatcagtgaa aaggtaaaca acaagggttaa tgtaactcta ttataaattt tgcatttttt      240
ttctctgtga catatacaag tatatttttg tttttggagc tataaattat ttaatttagc      300
aatcttcaaa gctcataaat ttcaactttt caaataagaa attttaactt caaataagaa      360
gtctaggact ttatggctat taattttact atcaaaatat ccaagggact ccattcaatg      420
taatagttat aattcttcta aatatcattt gaataattct ttgtggacgc tagactcaag      480
actatgctac atccaaacag tacatctata acc

```

<210> 315

<211> 222

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(222)

<223> n = A,T,C or G

<400> 315

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| atttatattc | aaggngatctc | aaagaaagca | ttttcatttc | actgcacatc | tagagaaaaa | 60 |
| caaaaataga | aaatcttcta | gtccatccta | atctgaatgg | tgctgtttct | atattgggtca | 120 |
| ttgccttgca | aacaggagct | ccacaaaagc | caggaagaga | gactgcctcc | ttggctgaaa | 180 |
| gagtcctttc | aggaaggtgg | actgcattgg | tttgatatgt | tt | | 222 |

<210> 316

<211> 1633

<212> DNA

<213> Homo sapiens

<400> 316

| | | | | | | |
|-------------|------------|------------|-------------|-------------|------------|------|
| cgtggaggca | gctagcgcga | ggctggggag | cgctgagccg | cgcgctcgtgc | cctgcgctgc | 60 |
| ccagactagc | gaacaatata | gtcgggatgg | ctaaagggtga | ccccaagaaa | ccaaagggca | 120 |
| agacgtccgc | ttatgccttc | tttgtgcaga | catgcagaga | agaacataag | aagaaaaacc | 180 |
| cagaggtccc | tgtcaatttt | gcggaatttt | ccaagaagtg | ctctgagagg | tggaagacgg | 240 |
| tgtccgggaa | agagaaatcc | aaatctgatg | aaatggcaaa | ggcagataaa | gtgcgctatg | 300 |
| atcgggaaat | gaaggattat | ggaccagcta | agggaggcaa | gaagaagaag | gatcctaata | 360 |
| ctcccaaaag | gccaccgtct | ggattcttcc | tggtctgttc | agaattccgc | cccaagatca | 420 |
| aatccacaaa | ccccggcatc | tctattggag | acgtggcaaa | aaagctgggt | gagatgtgga | 480 |
| ataatttaaa | tgacagtga | aagcagcctt | acatcactaa | ggcggcaaa | ctgaaggaga | 540 |
| agtatgagaa | ggatgttgct | gactataagt | cgaaaggaaa | gtttgatggg | gcaaaggggc | 600 |
| ctgctaaagt | tgcccgga | aaggtggaag | aggaagatga | agaacaggag | gaggaagaag | 660 |
| aggaggagga | ggaggaggag | gatgaataaa | gaaactgttt | atctgtctcc | ttgtgaatac | 720 |
| ttagagtagg | ggagcgccgt | aattgacaca | tctcttattt | gagaagtgtc | tggtgccctc | 780 |
| attaggttta | attacaaaat | ttgatcacga | tcattattgt | gtctctcaaa | gtgctctaga | 840 |
| aattgtcagt | ggtttacatg | aagtggccat | gggtgtctgg | agcacctga | aactgtatca | 900 |
| aagtgtgata | tatttccaaa | catttttaaa | atgaaaaggc | actctcgtgt | tctcctcact | 960 |
| ctgtgcactt | tgctgttggt | gtgacaaggc | atttaaagat | gtttctggca | ttttcttttt | 1020 |
| atgtgtaagg | tggtggtaac | tatggttatt | ggctagaaat | cctgagtttt | caactgtata | 1080 |
| tatctatagt | ttgtaaaaag | aacaaaacaa | ccgagacaaa | cccttgatgc | tccttgctcg | 1140 |
| gcgttgaggc | tgtggggaag | atgccttttg | ggagaggctg | tagctcaggg | cgtgcactgt | 1200 |
| gaggctggac | ctgttgactc | tgcaaggggc | atccatttag | cttcagggtg | tcttgcttct | 1260 |
| gtatatagt | acatagcatt | ctgctgccat | cttagctgtg | gacaaagggg | ggtcagctgg | 1320 |
| catgagaata | ttttttttta | agtgcggtag | tttttaaaact | gtttgttttt | aaacaaacta | 1380 |
| tagaactctt | cattgtcagc | aaagcaaaga | gtcactgcac | caatgaaagt | tcaagaacct | 1440 |
| cctgtactta | aacacgattc | gcaacgttct | gttatttttt | ttgtatgttt | agaatgctga | 1500 |
| aatgtttttg | aagttaaata | aacagtatta | cattttttaga | actcttctct | actataacag | 1560 |
| tcaattttctg | actcacagca | gtgaacaaac | ccccactccg | ttgtatttgg | agactggcct | 1620 |
| ccctataaat | gtg | | | | | 1633 |

<210> 317

<211> 4235

<212> DNA

<213> Homo sapiens

<400> 317

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| gaatccaagg | gggccagttc | ctgccgtctg | ctcttctgcc | tcttgatctc | cgccaccgtc | 60 |
| ttcaggccag | gccttggtatg | gtatactgta | aattcagcat | atggagatac | cattatcata | 120 |
| ccttgccgac | ttgacgtacc | tcagaatctc | atgtttggca | aatggaaata | tgaaaagccc | 180 |
| gatggctccc | cagtatttat | tgcttccaga | tcctctacaa | agaaaagtgt | gcagtacgac | 240 |
| gatgtaccag | aatacaaaga | cagattgaac | ctctcagaaa | actacacttt | gtctatcagt | 300 |
| aatgcaagga | tcagtgatga | aaagagattt | gtgtgcatgc | tagtaactga | ggacaacgtg | 360 |

tttgaggcac ctacaatagt caaggtgttc aagcaaccat ctaaacctga aattgtaagc 420
 aaagcactgt ttctcgaaac agagcagcta aaaaagttgg gtgactgcat ttcagaagac 480
 agttatccag atggcaatat cacatggtac aggaatggaa aagtgctaca tccccttgaa 540
 ggagcgggtg tcataatfff taaaaaggaa atggacccag tgactcagct ctataccatg 600
 acttccaccc tggagtacaa gacaaccaag gctgacatac aaatgccatt cactgtctcg 660
 gtgacatatt atggaccatc tggccagaaa acaattcatt ctgaacaggc agtatttgat 720
 atttactatc ctacagagca ggtgacaata caagtgtctgc caccacaaaa tgccatcaaa 780
 gaaggggata acatcactct taaatgctta gggaatggca accctcccc agaggaattt 840
 ttgttttact taccaggaca gcccgaagga ataagaagct caaatactta cactgtacg 900
 gatgtgaggc gcaatgcaac aggagactac aagtgttccc tgatagacaa aaaaagcatg 960
 attgttcaa cagccatcac agttcactat ttggatttgg ccttaaacc aagtggagaa 1020
 gtgactagac agattggtga tgccctaccc gtgtcatgca caatatctgc tagcaggaat 1080
 gcaactgtgg tatggatgaa agataacatc aggcctcgat ctagcccgctc attttctagt 1140
 cttcattatc aggatgctgg aaactatgtc tgcgaaactg ctctgcagga ggttgaagga 1200
 ctaaagaaaa gagagtcatt gactctcatt gtagaaggca aacctcaaat aaaaatgaca 1260
 aagaaaactg atcccagtg actatctaaa acaataatct gccatgtgga aggttttcca 1320
 aagccagcca ttcagtggac aattactggc agtggagcg tcataaacca aacagaggaa 1380
 tctccttata ttaatggcag gtattatagt aaaattatca tttccctga agagaatgtt 1440
 acattaactt gcacagcaga aaaccaactg gagagaacag taaactcctt gaatgtctct 1500
 gctataagta ttccagaaca cgatgaggca gacgagataa gtgatgaaa cagagaaaag 1560
 gtgaatgacc aggcaaaact aattgtggga atcgttgttg gtctcctcct tgctgccctt 1620
 gttgctggtg tctgtactg gctgtacatg aagaagtcaa agactgcac aaaacatgta 1680
 aacaaggacc tggtaatat ggaagaaaac aaaaagttag aagaaaacaa tcacaaaact 1740
 gaagcctaag agagaaaactg tcctagtgtt ccagagataa aatcatata gaccaattga 1800
 agcatgaacg tggattgtat ttaagacata aacaaagaca ttgacagcaa ttcattgtta 1860
 agtattaagc agttcattct accaagctgt cacaggtttt cagagaatta tctcaagtaa 1920
 aacaaatgaa atttaattac aaacaataag aacaagtttt ggcagccatg ataattggtc 1980
 atatgttgtg tttggttcaa ttttttttcc gtaaatgtct ctaattttat acactgtaag ctttgttctg 2040
 gtttgccctt tatgtaaatt ttttacgtga ctatttttat acactgtaag ctttgttctg 2100
 ggagttgctg ttaattctgat gtataatgta atgtttttat ttcaattgtt tatatggata 2160
 atctgagcag gtacatttct gattctgatt gctatcagca atgccccaaa ctttctcata 2220
 agcacctaaa acccaaaggt ggcagcttgt gaagattggg gacactcata ttgccctaat 2280
 taaaaactgt gatttttatc acaaggagg ggaggccgag agtcagactg atagacacca 2340
 taggagccga ctctttgata tgccaccagc gaactctcag aaataaatca cagatgcata 2400
 tagacacaca tacataatgg tactcccaaa ctgacaattt tacctattct gaaaaagaca 2460
 taaaacagaa tttggtagca cttacctcta cagacacctg ctaataaatt attttctgtc 2520
 aaaagaaaaa acacaagcat gtgtgagaga cagtttgga aaatcatggt caacattccc 2580
 attttcatag atcacaatgt aaatcactat aattacaaat tgggtgttaa tcctttgggt 2640
 tatccactgc cttaaaatta tacctatttc atgttttaaa agatatcaat cagaattgga 2700
 gtttttaaca gtggtcatta tcaaagctgt gttattttcc acagaatata gaatatatat 2760
 ttttttctg tgtgtttttg ttaactaccc tacagatatt gaatgcacct tgagataatt 2820
 tagtggttta actgatacat aatttatcaa gcagtacatg aaagtgtaat aataaaatgt 2880
 ctatgtatct ttagttacat tcaaatttgt aactttataa acatgtttta tgcttgagga 2940
 aatttttaag gtggtagtat aaatggaaac tttttgaagt agaccagata tgggctactt 3000
 gtgactagac ttttaaaact tgctctttca agcagaagcc tggtttctgg gagaacactg 3060
 cacagtgatt ttttccag gatttacaca actttaagg gaagataaat gaacatcaga 3120
 tttctaggta tagaactatg ttattgaaag gaaaaggaaa actggtgttt gtttcttaga 3180
 ctcatgaaat aaaaaattat gaaggcaatg aaaaataaat tgaaaattaa agtcagatga 3240
 gaataggaat aatactttgc cacttctgca ttatttagaa acatacgtaa ttgtacattt 3300
 gtaaacatt tactgtctgg gcaatagtga ctccgtttaa taaaagcttc cgtagtgcac 3360
 tggtaggat taaatgcata aaatatctta gactcgatgc tgtataaaat attatgggaa 3420
 aaaagaaata cgttattttg cctctaaact tttattgaag ttttatttgg caggaaaaaa 3480
 aattgaatct tggtaacat ttaaaccaaa gtaaaagggg aaaaaccaa gttatttgtt 3540
 ttgcatggct aagccattct gttatctctg taaatactgt gatttctttt ttattttctc 3600


```
<210> 319
<211> 1814
<212> DNA
<213> Homo sapiens
```

| | | | | | | | |
|------------|------------|-------------|-------------|--------------|-------------|------|--|
| <400> | 319 | | | | | | |
| ggggagatga | tccgagccgc | gccgcgcgcg | ctgttctctgc | tgtctgtctgct | gctgtctgctg | 60 | |
| ctagtgtcct | gggcgtccc | aggcgaggca | gcccccgacc | aggacgagat | ccagcgcctc | 120 | |
| cccgggctgg | ccaagcagcc | gtctttccgc | cagtactccg | gctacctcaa | aagctccggc | 180 | |
| tccaagcacc | tccactactg | gtttgtggag | tcccagaagg | atcccagagaa | cagccctgtg | 240 | |
| gtgctttggc | tcaatggggg | tcccggctgc | agctcactag | atgggctcct | cacagagcat | 300 | |
| ggcccccttc | tggtcacagc | agatgggtgc | accctggagt | acaaccctta | ttcttggaat | 360 | |
| ctgattgcc | atgtgttata | cctggagtc | ccagctgggg | tgggtttctc | ctactccgat | 420 | |
| gacaagtttt | atgcaactaa | tgacactgag | gtcgcccaga | gcaattttga | ggcccttcaa | 480 | |
| gatttcttcc | gcctcttttc | ggagtacaag | aacaacaaac | ttttctgac | cggggagagc | 540 | |
| tatgctggca | tctacatccc | caccttgccc | gtgctggcca | tgcaggatcc | cagcatgaac | 600 | |
| cttcaggggc | tggctgtggg | caatggagtc | tctctctatg | agcagaatga | caactccctg | 660 | |
| gtctactttg | cctactacca | tggccttctg | gggaacaggc | tttgggtctc | tctccagacc | 720 | |
| cactgctgct | ctcaaaaaca | gtgtaacttc | tatgacaaca | aagacctgga | atgctgacc | 780 | |
| aatcttcagg | aagtggcccg | catcgtgggc | aactctggcc | tcaacatcta | caatctctat | 840 | |
| gccccgtgtg | ctggaggggt | gccagccat | tttaggtatg | agaaggacac | tgttgtggtc | 900 | |
| caggatttgg | gcaacatctt | cactcgctg | ccactcaagc | ggatgtggca | tcaggcactg | 960 | |
| ctgcgctcag | gggataaagt | gcgcatggac | ccccctgca | ccaacacaac | agctgcttcc | 1020 | |
| acctacctca | acaaccctga | ctgctgggaag | gccctcaaca | tcccggagca | gctgccacaa | 1080 | |
| tgggacatgt | gcaactttct | ggtaaaactta | cagtaccgcc | gtctctaccg | aagcatgaac | 1140 | |
| tcccagtatc | tgaagctgct | tagctcacag | aaataccaga | tctattata | taatggagat | 1200 | |
| gtagacatgg | cctgcaattt | catgggggat | gagtggtttg | tggattccct | caaccagaag | 1260 | |
| atggaggtgc | agcgcgggcc | ctggttagtg | aagtaagggg | acagggggga | gcagattggc | 1320 | |
| ggcttcgtga | aggagtcttc | ccacatcgcc | tttctcacga | tcaagggcgc | cggccacatg | 1380 | |
| gttcccaccg | acaagccctc | cctgccttc | accatgttct | ccgccttctc | gaacaagcag | 1440 | |
| ccatactgat | gaccacagca | cagcgtcca | cggcctgatg | cagccccctc | cagcctctcc | 1500 | |
| cgctaggaga | gtcctcttct | aagcaaagtg | cccctgcagg | cggttctctg | cgccaggact | 1560 | |
| qcccccttcc | cagagccctg | tacatcccag | actgggcccc | gggtctccca | tagacagcct | 1620 | |

```

gggggcaagt tagcacttta ttcccgcagc agttcctgaa tgggggtggcc tggcccccttc 1680
tctgcttaaa gaatgccctt tatgatgcac tgattccatc ccaggaaccc aacagagctc 1740
aggacagccc acagggaggt ggtggacgga ctgtaattga tagattgatt atggaattaa 1800
attgggtaca gctt 1814

```

```

<210> 320
<211> 3132
<212> DNA
<213> Homo sapiens

```

```

<400> 320
ccgcagaact tggggagccg ccgcgcgcac ccgcgcgcgc agccagcttc cgccgcgcgc 60
ggaccggccc ctgccccagc ctccgcagcc gcggcgcgct caccgcccgc cgccgcgcagg 120
gcgagtcggg gtcgcgcgct gcacgcttct cagtgttccc cgccgcccgc atgtaacccg 180
gccaggcccc cgcaacgggtg tcccctgcag ctccagcccc gggctgcacc ccccgcccc 240
gacaccagct ctccagcctg ctcgccaggg atggccgcgg ccaaggccga gatgcagctg 300
atgtccccgc tgcagatctc tgaccgcttc ggatccttct ctcactcgcc caccatggac 360
aactacccta agctggagga gatgatgctg ctgagcaacg gggctcccca gttcctcggc 420
gccgcggggg ccccagaggg cagcggcagc aacagcagca gcagcagcag cgggggagggt 480
ggaggcggcg ggggcggcag caacagcagc agcagcagca gcaccttcaa ccctcaggcg 540
gacacggggc agcagcccta cgagcacctg accgcagagt cttttcctga catctctctg 600
aacaacgaga aggtgctggt ggagaccagt taccaccagc aaaccaactc actgcccccc 660
atcacctata ctggccgctt ttccctggag cctgcaccca acagtggcaa caccttgttg 720
cccagacccc tcttcagctt ggtcagtggc ctagtgcaga tgaccaaccc accggcctcc 780
tcgtcctcag caccatctcc agcggcctcc tccgcctccg cctcccagag cccacccctg 840
agctgcgcag tgccatccaa cgacagcagt cccatttact cagcggcacc caccttcccc 900
acgccgaaca ctgacatttt ccctgagcca caaagccagg ccttcccggg ctccggcagg 960
acagcgtccc agtaccgccc tctgcctac cctgcccaga aggggtggct ccagggtccc 1020
atgatccccg actacctgtt tccacagcag caggggggat tgggcctggg caccacagac 1080
cagaagccct tccagggcct ggagagccgc acccagcagc cttcgctaac cctctgtct 1140
actattaagg cctttgccac tcagtccggc tcccaggacc tgaaggccct caataaccag 1200
taccagtccc agctcatcaa acccagccgc atgcgcaagt atcccaaccg gccagcaag 1260
acgccccccc acgaacgccc ttacgcttgc ccagtggagt cctgtgatcg ccgcttctcc 1320
cgctccgacg agctcaccgg ccacatccgc atccacacag gccagaagcc cttccagtgc 1380
cgcatctgca tgcgcaactt cagccgcagc gaccacctca ccaccacat ccgcacccac 1440
acaggcgaaa agcccttcgc ctgcgacatc tgtggaagaa agtttgccag gagcgatgaa 1500
cgcaagaggc ataccaagat ccacttgccg cagaaggaca agaaagcaga caaaagtgtt 1560
gtggcctctt cggccacctc ctctctctct tcttaccgt ccccggttgc tacctcttac 1620
ccgtccccgg ttactacctc ttatccatcc ccggccacca cctcataccc atccccctgt 1680
cccacctcct tctctctctc cggctcctcg acctaccat cccctgtgca cagtggcttc 1740
ccctccccgt cgggtggccac cagctactcc tctgttcccc ctgctttccc ggcccagggt 1800
agcagcttcc cttcctcagc tgtaccaac tcttcagcg cctccacagg gctttcggac 1860
atgacagcaa ccttttctcc caggacaatt gaaatttgct aaagggaag gggaaagaaa 1920
gggaaaaggg agaaaaagaa acacaagaga cttaaaggac aggaggagga gatggccata 1980
ggagaggagg gttcctctta ggtcagatgg aggttctcag agccaagtcc tccctctcta 2040
ctggagtggg aggtctattg gccacaatc ctttctgccc acttcccctt cccaattac 2100
tattcccttt gacttcagct gcctgaaaca gccatgtcca agttcttcac ctctatccaa 2160
agaacttgat ttgcatggat tttggataaa tcatttcagt atcatctcca tcatatgcct 2220
gaccccttgc tcccttcaat gctagaaaat cgagttggca aaatgggggt tgggcccctc 2280
agagccctgc cctgcaccct tgtacagtgt ctgtgccatg gatttcgttt ttcttggggt 2340
actcttgatg tgaagataat ttgcatattc tattgtatta tttggagtta ggtcctcact 2400
tgggggaaaa aaaaaaaaaa aagccaagca aaccaatggg gatcctctat tttgtgatga 2460
tgctgtgaca ataagtttga accttttttt ttgaaacagc agtcccagta ttctcagagc 2520
atgtgtcaga gtgttgttcc gttaaccttt ttgtaaatat tgcttgaccg tactctcaca 2580

```

```

tgtggcaaaa tatggttttg tttttctttt ttttttttga aagtgttttt tcttcgtcct 2640
tttggtttta aaagtttcac gtcttggtgc cttttgtgtg atgccccttg ctgatggctt 2700
gacatgtgca attgtgaggg acatgctcac ctctagcctt aaggggggca gggagtgatg 2760
atttggggga ggctttggga gcaaaaataag gaagagggtc gagctgagct tcggttctcc 2820
agaatgtaag aaaacaaaat ctaaaacaaa atctgaactc tcaaaagtct atttttttaa 2880
ctgaaaatgt aaattttataa atatattcag gagttggaat gttgtagtta cctactgagt 2940
aggcggcgat ttttgtatgt tatgaacatg cagttcatta ttttgtgggt ctattttact 3000
ttgtacttgt gtttgcctaa acaaagtgcac tgtttggctt ataaacacat tgaatgcgct 3060
ttattgcccc tgggatatgt ggtgtatata cttccaaaaa attaaaacga aaataaagta 3120
gctgcgattg gg 3132

```

<210> 321

<211> 2280

<212> DNA

<213> Homo sapiens

<400> 321

```

ccgcccgcga ccagctacgc cccgtccgac gtgccctcgg gggtcgcgct gttcctcacc 60
atccctttcg ccttcttcct gcccgagctg atattttgggt tcttgggtctg gaccatggta 120
gccgccaccc acatagtata ccccttgctg caaggatggg tgatgtatgt ctgcctcacc 180
tcgtttctca tctccttgat gttcctggtt tcttacttgt ttggatttta caaaagattt 240
gaatcctgga gagttctgga cagcctgtac cacgggacca ctggcatcct gtacatgagc 300
gctgcctgcc tacaagtaca tgccacgatt gtttctgaga aactgctgga cccaagaatt 360
tactacatta attcggcagc ctgcgttcttc gccttcacatg ccacgctgct ctacattctc 420
catgccttca gcacttatta ccactgatgc acaggcgcca ggccaagggg gaaatgctct 480
ttgaaagctc caattatttg tccccaaaag cagcttccaa cgtttgccat ctggatgaca 540
aacggaagat ccactaaaac gtccacggga ttaacagaac gtccttgacg actgagcgat 600
gacaccacac tttgtttgga catttaaatt cactctgctg aataggagga agcttttctt 660
tttctgggga aaacaactgt ctcttggaat tatctgacca tgaacttgct cttctagaca 720
actcacatca aagccctcac tccactaatg gagaatccta gccccactaa tgccaagtct 780
gtttggggat tttgcctcag ctatgggctt ccctagagta ggtctagggg aatactcagt 840
ctgatctttt ttttgtttgt tttattttgt tttttttgag acggagtctc gctcttctct 900
caaggctgga gtgcagtgc gcgatctcca ctactgcag gctccgcctc ccgggttccc 960
gccattctcc tgctcagcc tcccgagtag ccgggactac aggcgcccac caccatgccc 1020
ggctaattta gttgtatttt tagtagagat ggggtttcac cgtattagcc aggatggct 1080
cgatctcctg acctcgtgat ccgcccgcct cggctcccca aagtgcctggg attacaggcg 1140
tgagccaccg tgcccggcct gattctctta aaattgaaga ggtgctgcca aggccctcag 1200
atctaacgca gatgcataga cctgttctct ggtacttggt cagcctgtgc tggggagccg 1260
tggtcccagag ttccctggga ggctgacagg gtcaagccac cctgcccacc accctcccac 1320
ttccctctcc ctttctctc cagcattagg attcaaggga aatctgcatg aagccaattt 1380
tgagggtaga cgtgtgggga aaataaatca ttatacagta agacctgggg cttgaggggt 1440
ggggaatggg gaggggaagg catagcctgc tctccatga gtctgacatc tcggaaactg 1500
agcagctgcc ggacgcctgg gtcaggaatc caagaccca cctcttaagg actggttcct 1560
cagaaagcac cctcagggaa aaagggtgaaa acattacatc cgtggattct cctgccacaa 1620
ccgcatgtga agaaaaggc gccgcaacat ctgagcgagg agtgaaggac ccatgtccca 1680
ggaaccgcgc tgcgccacct gcactacccc ccctcacatt ctcttaagca cccggtggcc 1740
ctccgaggct ggcggaatgg tgggtgccac ggggttgggc aagggtcac caggacctca 1800
acgggcaaaag ttgtgcacac taaaatatca aatcaagggt cttggtttta aagtaaattg 1860
ttttctaaag aaagctgtgt tcttctgttg acccagacga atagggcaca gccctgtaac 1920
tgacagtgcc ttctgtcatt gggaatgaaa taaattatta cgagaaaggg acttgtccta 1980
actggtttga ggccttacag ttttgtatct acatttttcc cctcctgggg tttgcgggga 2040
cagggacaga actacaggag tcatgggaaa gaaaattctg gcttcactac tgctcactgc 2100
tcactttctg atcactctga tacttttttt tttttttttt ttttgcaacc tgataccttg 2160
aaaagcttct atgtgtctct ccttttgttg cctggcagct gtctaggatg atcactgatt 2220

```

actattttact aagtagccac atgcaaataa aagttgtttg gtaaaatgga aaaaaaaaaa 2280

<210> 322

<211> 1398

<212> DNA

<213> Homo sapiens

<400> 322

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|------------|------|
| tagatggcaa | cctccctatc | tgcccgagg | tcatagaggc | gacacgtagc | gtcatctgac | 60 |
| cctgaagcaa | aggcatctcc | actccaaagt | tagacaaaat | gccaggaatg | ttcttctctg | 120 |
| ctaaccctaaa | ggaattgaaa | ggaaccactc | attcacttct | agacgacaaa | atgcaaaaaa | 180 |
| ggaggccaaa | gacttttgga | atggatatga | aagcatacct | gagatctatg | atcccacatc | 240 |
| tggaatctgg | aatgaaatct | tccaagtcca | aggatgtact | ttctgctgct | gaagtaatgc | 300 |
| aatggtctca | atctctggaa | aaacttcttg | ccaacccaaac | tgggtcaaat | gtctttggaa | 360 |
| gtttcctaaa | gtctgaattc | agtgaggaga | atattgagtt | ctggctggct | tgtgaagact | 420 |
| ataagaaaac | agagtctgat | cttttgcctt | gtaaagcaga | agagatatat | aaagcatttg | 480 |
| tgcattcaga | tgtctgctaaa | caaatacaata | ttgacttccg | cactcgagaa | tctacagcca | 540 |
| agaagattaa | agcaccaacc | cccacgtgtt | ttgatgaagc | acaaaaagtc | atatatactc | 600 |
| ttatggaaaa | ggactcttat | cccagggttc | tcaaatacaga | tatttactta | aatcttctaa | 660 |
| atgacctgca | ggctaatagc | ctaaagtgac | tgggtccctgg | ctgaagggaa | ttaacagata | 720 |
| gtatcaaggc | acgaaggaa | gtgccagtat | ggctccctgg | gtgaacagct | tggccttttt | 780 |
| tgggtgtctt | gacaggccaa | gaagaacaaa | tgactcagaa | tggattaaca | tgaaagttat | 840 |
| ccaggcgagc | agttgaagaa | gcataagcaa | gacaaaaaca | gagagaccgc | agaaggagga | 900 |
| agatactgtg | gtactgtcat | aaaaaacagt | ggagctctgt | attagaaagc | ccctcagaac | 960 |
| tgggaaggcc | aggtaactct | agttacacag | aaactgtgac | taaagtctat | gaaactgatt | 1020 |
| acaacaggct | gtaagaatca | aagtcaactg | acatctatgc | tacatatatt | tatatagttt | 1080 |
| gtactgagct | attgaagtcc | cattaactta | aagtatatgt | tttcaaattg | ccattgctac | 1140 |
| tattgcttgt | cgggtgattt | tattttattg | tttttgactt | tgggaagagat | gaactgtgta | 1200 |
| tttaacttaa | gctattgtct | ttaaaaccag | ggatcagaat | atattttgtaa | gttaaatcat | 1260 |
| tgggtgcta | aatgaaatgtg | gattttgtat | taaaatatat | agaagcaatt | tctgtttaca | 1320 |
| tgtccttgct | acttttaaaa | acttgcattt | attcctcaga | ttttaaaaat | aaataaataa | 1380 |
| ttcattttaa | aaaaaaaa | | | | | 1398 |

<210> 323

<211> 1316

<212> DNA

<213> Homo sapiens

<400> 323

| | | | | | | |
|-------------|-------------|------------|------------|-------------|------------|-----|
| acttctacct | gtcactcag | aatcatttct | gcaccaacca | tggccacgtt | tgtggagctc | 60 |
| agtaccaaag | ccaagatgcc | cattgtgggc | ctgggcactt | ggaagtctcc | tcttggcaaa | 120 |
| gtgaaagaag | cagtgaaggt | ggccattgat | gcaggatata | ggcacattga | ctgtgcctat | 180 |
| gtctatcaga | atgaacatga | agtgggggaa | gccatccaag | agaagatcca | agagaaggct | 240 |
| gtgaagcggg | aggacctgtt | catcgtcagc | aagttgtggc | ccactttctt | tgagagaccc | 300 |
| cttgtgagga | aagcctttga | gaagaccctc | aaggacctga | agctgagcta | tctggacgtc | 360 |
| tattcttattc | actggccaca | gggattcaag | tctggggatg | accttttccc | caaagatgat | 420 |
| aaaggtaatg | ccatcggtgg | aaaagcaacg | ttcttggatg | cctggggaggc | catggaggag | 480 |
| ctggtggatg | aggggctgg | gaaagccctt | ggggtctcca | atttcagcca | cttccagatc | 540 |
| gagaagctct | tgaacaaacc | tggactgaaa | tataaaccag | tgactaacca | ggttgagtgt | 600 |
| caccataacc | tcacacagga | gaaactgata | cagtactgcc | actccaaggg | catcaccgtt | 660 |
| acggcctaca | gccccctggg | ctctccggat | agaccttggg | ccaagccaga | agacccttcc | 720 |
| ctgctggagg | atoccaaagat | taaggagatt | gctgcaaagc | acaaaaaac | cgcagcccag | 780 |
| gttctgatcc | gtttccatat | ccagaggaat | gtgattgtca | tccccaagtc | tgtgacacca | 840 |
| gcacgcattg | ttgagaacat | tcagggtctt | gacttttaaa | tgagtgatga | ggagatggca | 900 |

accatactca gcttcaacag aaactggagg gcctgtaacg tgttgcaatc ctctcatttg 960
gaagactatc ccttcaatgc agaattattga ggttgaatct cctgggtgaga ttatacagga 1020
gattctcttt ctctgctgaa gtgtgactac ctccactcat gtcccathtt agccaagctt 1080
atttaagatc acagtgaact tagtcctgtt atagacgaga atcgaggtgc tgttttagac 1140
atttatttct gtatgttcaa ctaggatcag aatatcacag aaaagcatgg cttgaataag 1200
gaaatgacaa ttttttccac ttatctgac agaacaaatg tttattaagc atcagaaact 1260
ctgccaacac tgaggatgta aagatcaata aaacaaataa taatcataaa aaaaaa 1316

<210> 324

<211> 200

<212> PRT

<213> Homo sapiens

<400> 324

Met Ala Lys Gly Asp Pro Lys Lys Pro Lys Gly Lys Thr Ser Ala Tyr
5 10 15

Ala Phe Phe Val Gln Thr Cys Arg Glu Glu His Lys Lys Lys Asn Pro
20 25 30

Glu Val Pro Val Asn Phe Ala Glu Phe Ser Lys Lys Cys Ser Glu Arg
35 40 45

Trp Lys Thr Val Ser Gly Lys Glu Lys Ser Lys Phe Asp Glu Met Ala
50 55 60

Lys Ala Asp Lys Val Arg Tyr Asp Arg Glu Met Lys Asp Tyr Gly Pro
65 70 75 80

Ala Lys Gly Gly Lys Lys Lys Lys Asp Pro Asn Ala Pro Lys Arg Pro
85 90 95

Pro Ser Gly Phe Phe Leu Phe Cys Ser Glu Phe Arg Pro Lys Ile Lys
100 105 110

Ser Thr Asn Pro Gly Ile Ser Ile Gly Asp Val Ala Lys Lys Leu Gly
115 120 125

Glu Met Trp Asn Asn Leu Asn Asp Ser Glu Lys Gln Pro Tyr Ile Thr
130 135 140

Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp Val Ala Asp Tyr
145 150 155 160

Lys Ser Lys Gly Lys Phe Asp Gly Ala Lys Gly Pro Ala Lys Val Ala
165 170 175

Arg Lys Lys Val Glu Glu Glu Asp Glu Glu Gln Glu Glu Glu Glu Glu
180 185 190

Glu Glu Glu Glu Glu Glu Asp Glu
195 200

006230" 036F3360

<211> 263

<212> PRT

<213> Homo sapiens

<400> 325

Met Phe Arg Asn Gln Tyr Asp Asn Asp Val Thr Val Trp Ser Pro Gln
5 10 15

Gly Arg Ile His Gln Ile Glu Tyr Ala Met Glu Ala Val Lys Gln Gly
20 25 30

Ser Ala Thr Val Gly Leu Lys Ser Lys Thr His Ala Val Leu Val Ala
35 40 45

Leu Lys Arg Ala Gln Ser Glu Leu Ala Ala His Gln Lys Lys Ile Leu
50 55 60

His Val Asp Asn His Ile Gly Ile Ser Ile Ala Gly Leu Thr Ala Asp
65 70 75 80

Ala Arg Leu Leu Cys Asn Phe Met Arg Gln Glu Cys Leu Asp Ser Arg
85 90 95

Phe Val Phe Asp Arg Pro Leu Pro Val Ser Arg Leu Val Ser Leu Ile
100 105 110

Gly Ser Lys Thr Gln Ile Pro Thr Gln Arg Tyr Gly Arg Arg Pro Tyr
115 120 125

Gly Val Gly Leu Leu Ile Ala Gly Tyr Asp Asp Met Gly Pro His Ile
130 135 140

Phe Gln Thr Cys Pro Ser Ala Asn Tyr Phe Asp Cys Arg Ala Met Ser
145 150 155 160

Ile Gly Ala Arg Ser Gln Ser Ala Arg Thr Tyr Leu Glu Arg His Met
165 170 175

Ser Glu Phe Met Glu Cys Asn Leu Asn Glu Leu Val Lys His Gly Leu
180 185 190

Arg Ala Leu Arg Glu Thr Leu Pro Ala Glu Gln Asp Leu Thr Thr Lys
195 200 205

Asn Val Ser Ile Gly Ile Val Gly Lys Asp Leu Glu Phe Thr Ile Tyr
210 215 220

Asp Asp Asp Asp Val Ser Pro Phe Leu Glu Gly Leu Glu Glu Arg Pro
225 230 235 240

Gln Arg Lys Ala Gln Pro Ala Gln Pro Ala Asp Glu Pro Ala Glu Lys
245 250 255

<210> 326

<211> 539

<212> PRT

<213> Homo sapiens

<400> 326

Met Pro Glu Asn Val Ala Pro Arg Ser Gly Ala Thr Ala Gly Ala Ala

Gly Gly Arg Gly Lys Gly Ala Tyr Gln Asp Arg Asp Lys Pro Ala Gln
20 25 30

Ile Arg Phe Ser Asn Ile Ser Ala Ala Lys Ala Val Ala Asp Ala Ile
35 40 45

Arg Thr Ser Leu Gly Pro Lys Gly Met Asp Lys Met Ile Gln Asp Gly
50 55 60

Lys Gly Asp Val Thr Ile Thr Asn Asp Gly Ala Thr Ile Leu Lys Gln
65 70 75 80

Met Gln Val Leu His Pro Ala Ala Arg Met Leu Val Glu Leu Ser Lys
85 90 95

Ala Gln Asp Ile Glu Ala Gly Asp Gly Thr Thr Ser Val Val Ile Ile
100 105 110

Ala Gly Ser Leu Leu Asp Ser Cys Thr Lys Leu Leu Gln Lys Gly Ile
115 120 125

His Pro Thr Ile Ile Ser Glu Ser Phe Gln Lys Ala Leu Glu Lys Gly
130 135 140

Ile Glu Ile Leu Thr Asp Met Ser Arg Pro Val Glu Leu Ser Asp Arg
145 150 155 160

Glu Thr Leu Leu Asn Ser Ala Thr Thr Ser Leu Asn Ser Lys Val Val
165 170 175

Ser Gln Tyr Ser Ser Leu Leu Ser Pro Met Ser Val Asn Ala Val Met
180 185 190

Lys Val Ile Asp Pro Ala Thr Ala Thr Ser Val Asp Leu Arg Asp Ile
195 200 205

Lys Ile Val Lys Lys Leu Gly Gly Thr Ile Asp Asp Cys Glu Leu Val
210 215 220

Glu Gly Leu Val Leu Thr Gln Lys Val Ser Asn Ser Gly Ile Thr Arg
225 230 235 240

Val Glu Lys Ala Lys Ile Gly Leu Ile Gln Phe Cys Leu Ser Ala Pro
 245 250 255
 Lys Thr Asp Met Asp Asn Gln Ile Val Val Ser Asp Tyr Ala Gln Met
 260 265 270
 Asp Arg Val Leu Arg Glu Glu Arg Ala Tyr Ile Leu Asn Leu Val Lys
 275 280 285
 Gln Ile Lys Lys Thr Gly Cys Asn Val Leu Leu Ile Gln Lys Ser Ile
 290 295 300
 Leu Arg Asp Ala Leu Ser Asp Leu Ala Leu His Phe Leu Asn Lys Met
 305 310 315 320
 Lys Ile Met Val Ile Lys Asp Ile Glu Arg Glu Asp Ile Glu Phe Ile
 325 330 335
 Cys Lys Thr Ile Gly Thr Lys Pro Val Ala His Ile Asp Gln Phe Thr
 340 345 350
 Ala Asp Met Leu Gly Ser Ala Glu Leu Ala Glu Glu Val Asn Leu Asn
 355 360 365
 Gly Ser Gly Lys Leu Leu Lys Ile Thr Gly Cys Ala Ser Pro Gly Lys
 370 375 380
 Thr Val Thr Ile Val Val Arg Gly Ser Asn Lys Leu Val Ile Glu Glu
 385 390 395 400
 Ala Glu Arg Ser Ile His Asp Ala Leu Cys Val Ile Arg Cys Leu Val
 405 410 415
 Lys Lys Arg Ala Leu Ile Ala Gly Gly Gly Ala Pro Glu Ile Glu Leu
 420 425 430
 Ala Leu Arg Leu Thr Glu Tyr Ser Arg Thr Leu Ser Gly Met Glu Ser
 435 440 445
 Tyr Cys Val Arg Ala Phe Ala Asp Ala Met Glu Val Ile Pro Ser Thr
 450 455 460
 Leu Ala Glu Asn Ala Gly Leu Asn Pro Ile Ser Thr Val Thr Glu Leu
 465 470 475 480
 Arg Asn Arg His Ala Gln Gly Glu Lys Thr Ala Gly Ile Asn Val Arg
 485 490 495
 Lys Gly Gly Ile Ser Asn Ile Leu Glu Glu Leu Val Val Gln Pro Leu
 500 505 510
 Leu Val Ser Val Ser Ala Leu Thr Leu Ala Thr Glu Thr Val Arg Ser
 515 520 525

```
<210> 327
<211> 144
<212> PRT
<213> Homo sapiens
```

Thr Ala Ala Leu Ile Phe Phe Ala Ile Trp His Ile Ile Ala Phe Asp
20 25 30

Asn Pro Leu Val Leu Pro Glu Tyr Leu Ile His Ala Phe Phe Cys Val
50 55 60

Leu Leu Ala Tyr His Ile Trp Arg Tyr Met Ser Arg Pro Val Met Ser
85 90 95

Ala Tyr Cys Gln Lys Glu Gly Trp Cys Lys Leu Ala Phe Tyr Leu Leu
115 120 125

```
<210> 328
<211> 138
<212> PRT
<213> Homo sapiens
```

Glu Glu Leu Leu Lys Val Leu Gly Val Asn Val Met Leu Arg Lys Ile
20 25 30

Ala Val Ala Ala Ala Ser Lys Pro Ala Val Glu Ile Lys Gln Glu Gly
35 40 45

Asp Leu Pro Glu Asp Val Lys Trp Ile Asp Ile Thr Pro Asp Met Met
145 150 155 160

Asn Val Ser Ser His Leu Asp Lys Ala Ser Val Met Arg Leu Thr Ile

340 345 350
 Ser Leu Gln Gln Thr Glu Cys Val Leu Lys Pro Val Glu Ser Ser Asp
 355 360 365
 Met Lys Met Thr Gln Leu Phe Thr Lys Val Glu Ser Glu Asp Thr Ser
 370 375 380
 Ser Leu Phe Asp Lys Leu Lys Lys Glu Pro Asp Ala Leu Thr Leu Leu
 385 390 395 400
 Ala Pro Ala Ala Gly Asp Thr Ile Ile Ser Leu Asp Phe Gly Ser Asn
 405 410 415
 Asp Thr Glu Thr Asp Asp Gln Gln Leu Glu Glu Val Pro Leu Tyr Asn
 420 425 430
 Asp Val Met Leu Pro Ser Pro Asn Glu Lys Leu Gln Asn Ile Asn Leu
 435 440 445
 Ala Met Ser Pro Leu Pro Thr Ala Glu Thr Pro Lys Pro Leu Arg Ser
 450 455 460
 Ser Ala Asp Pro Ala Leu Asn Gln Glu Val Ala Leu Lys Leu Glu Pro
 465 470 475 480
 Asn Pro Glu Ser Leu Glu Leu Ser Phe Thr Met Pro Gln Ile Gln Asp
 485 490 495
 Gln Thr Pro Ser Pro Ser Asp Gly Ser Thr Arg Gln Ser Ser Pro Glu
 500 505 510
 Pro Asn Ser Pro Ser Glu Tyr Cys Phe Tyr Val Asp Ser Asp Met Val
 515 520 525
 Asn Glu Phe Lys Leu Glu Leu Val Glu Lys Leu Phe Ala Glu Asp Thr
 530 535 540
 Glu Ala Lys Asn Pro Phe Ser Thr Gln Asp Thr Asp Leu Asp Leu Glu
 545 550 555 560
 Met Leu Ala Pro Tyr Ile Pro Met Asp Asp Asp Phe Gln Leu Arg Ser
 565 570 575
 Phe Asp Gln Leu Ser Pro Leu Glu Ser Ser Ser Ala Ser Pro Glu Ser
 580 585 590
 Ala Ser Pro Gln Ser Thr Val Thr Val Phe Gln Gln Thr Gln Ile Gln
 595 600 605
 Glu Pro Thr Ala Asn Ala Thr Thr Thr Thr Ala Thr Thr Asp Glu Leu
 610 615 620
 Lys Thr Val Thr Lys Asp Arg Met Glu Asp Ile Lys Ile Leu Ile Ala

```
<210> 331
<211> 92
<212> PRT
<213> Homo sapiens
```

```

<400> 331
Met Ala Tyr Arg Gly Gln Gly Gln Lys Val Gln Lys Val Met Val Gln
                    5                      10                      15

Pro Ile Asn Leu Ile Phe Arg Tyr Leu Gln Asn Arg Ser Arg Ile Gln
                20                      25                      30

Val Trp Leu Tyr Glu Gln Val Asn Met Arg Ile Glu Gly Cys Ile Ile
    35                      40                      45

```

His Ser Lys Thr Lys Ser Arg Lys Gln Leu Gly Arg Ile Met Leu Lys
65 70 75 80

<210> 332

<211> 235

<212> PRT

<213> Homo sapiens

<400> 332

Met Asp Pro Ala Arg Pro Leu Gly Leu Ser Ile Leu Leu Leu Phe Leu

Thr Glu Ala Ala Leu Gly Asp Ala Ala Gln Glu Pro Thr Gly Asn Asn
20 25 30

Ala Glu Ile Cys Leu Leu Pro Leu Asp Tyr Gly Pro Cys Arg Ala Leu
35 40 45

Leu Leu Arg Tyr Tyr Tyr Asp Arg Tyr Thr Gln Ser Cys Arg Gln Phe
50 55 60

Leu Tyr Gly Gly Cys Glu Gly Asn Ala Asn Asn Phe Tyr Thr Trp Glu
65 70 75 80

Ala Cys Asp Asp Ala Cys Trp Arg Ile Glu Lys Val Pro Lys Val Cys
85 90 95

Arg Leu Gln Val Ser Val Asp Asp Gln Cys Glu Gly Ser Thr Glu Lys
100 105 110

Tyr Phe Phe Asn Leu Ser Ser Met Thr Cys Glu Lys Phe Phe Ser Gly
115 120 125

Gly Cys His Arg Asn Arg Ile Glu Asn Arg Phe Pro Asp Glu Ala Thr
130 135 140

Cys Met Gly Phe Cys Ala Pro Lys Lys Ile Pro Ser Phe Cys Tyr Ser
145 150 155 160

Pro Lys Asp Glu Gly Leu Cys Ser Ala Asn Val Thr Arg Tyr Tyr Phe
165 170 175

Asn Pro Arg Tyr Arg Thr Cys Asp Ala Phe Thr Tyr Thr Gly Cys Gly
180 185 190

Gly Asn Asp Asn Asn Phe Val Ser Arg Glu Asp Cys Lys Arg Ala Cys
195 200 205

Ala Lys Ala Leu Lys Lys Lys Lys Lys Met Pro Lys Leu Arg Phe Ala
 210 215 220

Ser Arg Ile Arg Lys Ile Arg Lys Lys Gln Phe
 225 230 235

<210> 333
 <211> 291
 <212> PRT
 <213> Homo sapiens

<400> 333
 Met Gln Arg Ala Arg Pro Thr Leu Trp Ala Ala Ala Leu Thr Leu Leu
 5 10 15

Val Leu Leu Arg Gly Pro Pro Val Ala Arg Ala Gly Ala Ser Ser Gly
 20 25 30

Gly Leu Gly Pro Val Val Arg Cys Glu Pro Cys Asp Ala Arg Ala Leu
 35 40 45

Ala Gln Cys Ala Pro Pro Pro Ala Val Cys Ala Glu Leu Val Arg Glu
 50 55 60

Pro Gly Cys Gly Cys Cys Leu Thr Cys Ala Leu Ser Glu Gly Gln Pro
 65 70 75 80

Cys Gly Ile Tyr Thr Glu Arg Cys Gly Ser Gly Leu Arg Cys Gln Pro
 85 90 95

Ser Pro Asp Glu Ala Arg Pro Leu Gln Ala Leu Leu Asp Gly Arg Gly
 100 105 110

Leu Cys Val Asn Ala Ser Ala Val Ser Arg Leu Arg Ala Tyr Leu Leu
 115 120 125

Pro Ala Pro Pro Ala Pro Gly Asn Ala Ser Glu Ser Glu Glu Asp Arg
 130 135 140

Ser Ala Gly Ser Val Glu Ser Pro Ser Val Ser Ser Thr His Arg Val
 145 150 155 160

Ser Asp Pro Lys Phe His Pro Leu His Ser Lys Ile Ile Ile Ile Lys
 165 170 175

Lys Gly His Ala Lys Asp Ser Gln Arg Tyr Lys Val Asp Tyr Glu Ser
 180 185 190

Gln Ser Thr Asp Thr Gln Asn Phe Ser Ser Glu Ser Lys Arg Glu Thr
 195 200 205

Glu Tyr Gly Pro Cys Arg Arg Glu Met Glu Asp Thr Leu Asn His Leu
 210 215 220

006260" E35T360

Lys Phe Leu Asn Val Leu Ser Pro Arg Gly Val His Ile Pro Asn Cys
225 230 235 240

Asp Lys Lys Gly Phe Tyr Lys Lys Lys Gln Cys Arg Pro Ser Lys Gly
245 250 255

Arg Lys Arg Gly Phe Cys Trp Cys Val Asp Lys Tyr Gly Gln Pro Leu
260 265 270

Pro Gly Tyr Thr Thr Lys Gly Lys Glu Asp Val His Cys Tyr Ser Met
275 280 285

Gln Ser Lys
290

<210> 334

<211> 582

<212> PRT

<213> Homo sapiens

<400> 334

Glu Ser Lys Gly Ala Ser Ser Cys Arg Leu Leu Phe Cys Leu Leu Ile
5 10 15

Ser Ala Thr Val Phe Arg Pro Gly Leu Gly Trp Tyr Thr Val Asn Ser
20 25 30

Ala Tyr Gly Asp Thr Ile Ile Ile Pro Cys Arg Leu Asp Val Pro Gln
35 40 45

Asn Leu Met Phe Gly Lys Trp Lys Tyr Glu Lys Pro Asp Gly Ser Pro
50 55 60

Val Phe Ile Ala Phe Arg Ser Ser Thr Lys Lys Ser Val Gln Tyr Asp
65 70 75 80

Asp Val Pro Glu Tyr Lys Asp Arg Leu Asn Leu Ser Glu Asn Tyr Thr
85 90 95

Leu Ser Ile Ser Asn Ala Arg Ile Ser Asp Glu Lys Arg Phe Val Cys
100 105 110

Met Leu Val Thr Glu Asp Asn Val Phe Glu Ala Pro Thr Ile Val Lys
115 120 125

Val Phe Lys Gln Pro Ser Lys Pro Glu Ile Val Ser Lys Ala Leu Phe
130 135 140

Leu Glu Thr Glu Gln Leu Lys Lys Leu Gly Asp Cys Ile Ser Glu Asp
145 150 155 160

Ser Tyr Pro Asp Gly Asn Ile Thr Trp Tyr Arg Asn Gly Lys Val Leu

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | 165 | | 170 | | 175 | | | | | | | | | | |
| His | Pro | Leu | Glu | Gly | Ala | Val | Val | Ile | Ile | Phe | Lys | Lys | Glu | Met | Asp |
| | 180 | | | | | | | 185 | | | | | 190 | | |
| Pro | Val | Thr | Gln | Leu | Tyr | Thr | Met | Thr | Ser | Thr | Leu | Glu | Tyr | Lys | Thr |
| | 195 | | | | | | 200 | | | | | 205 | | | |
| Thr | Lys | Ala | Asp | Ile | Gln | Met | Pro | Phe | Thr | Cys | Ser | Val | Thr | Tyr | Tyr |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Gly | Pro | Ser | Gly | Gln | Lys | Thr | Ile | His | Ser | Glu | Gln | Ala | Val | Phe | Asp |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Ile | Tyr | Tyr | Pro | Thr | Glu | Gln | Val | Thr | Ile | Gln | Val | Leu | Pro | Pro | Lys |
| | | | | 245 | | | | | 250 | | | | | | 255 |
| Asn | Ala | Ile | Lys | Glu | Gly | Asp | Asn | Ile | Thr | Leu | Lys | Cys | Leu | Gly | Asn |
| | | | 260 | | | | | 265 | | | | | | 270 | |
| Gly | Asn | Pro | Pro | Pro | Glu | Glu | Phe | Leu | Phe | Tyr | Leu | Pro | Gly | Gln | Pro |
| | 275 | | | | | | 280 | | | | | 285 | | | |
| Glu | Gly | Ile | Arg | Ser | Ser | Asn | Thr | Tyr | Thr | Leu | Thr | Asp | Val | Arg | Arg |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Asn | Ala | Thr | Gly | Asp | Tyr | Lys | Cys | Ser | Leu | Ile | Asp | Lys | Lys | Ser | Met |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Ile | Ala | Ser | Thr | Ala | Ile | Thr | Val | His | Tyr | Leu | Asp | Leu | Ser | Leu | Asn |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Pro | Ser | Gly | Glu | Val | Thr | Arg | Gln | Ile | Gly | Asp | Ala | Leu | Pro | Val | Ser |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| Cys | Thr | Ile | Ser | Ala | Ser | Arg | Asn | Ala | Thr | Val | Val | Trp | Met | Lys | Asp |
| | | 355 | | | | | 360 | | | | | 365 | | | |
| Asn | Ile | Arg | Leu | Arg | Ser | Ser | Pro | Ser | Phe | Ser | Ser | Leu | His | Tyr | Gln |
| | 370 | | | | | 375 | | | | | 380 | | | | |
| Asp | Ala | Gly | Asn | Tyr | Val | Cys | Glu | Thr | Ala | Leu | Gln | Glu | Val | Glu | Gly |
| 385 | | | | | 390 | | | | | 395 | | | | | 400 |
| Leu | Lys | Lys | Arg | Glu | Ser | Leu | Thr | Leu | Ile | Val | Glu | Gly | Lys | Pro | Gln |
| | | | | 405 | | | | | 410 | | | | | 415 | |
| Ile | Lys | Met | Thr | Lys | Lys | Thr | Asp | Pro | Ser | Gly | Leu | Ser | Lys | Thr | Ile |
| | | | 420 | | | | | 425 | | | | | 430 | | |
| Ile | Cys | His | Val | Glu | Gly | Phe | Pro | Lys | Pro | Ala | Ile | Gln | Trp | Thr | Ile |
| | 435 | | | | | | 440 | | | | | 445 | | | |
| Thr | Gly | Ser | Gly | Ser | Val | Ile | Asn | Gln | Thr | Glu | Glu | Ser | Pro | Tyr | Ile |

```

450              455              460
Asn Gly Arg Tyr Tyr Ser Lys Ile Ile Ile Ser Pro Glu Glu Asn Val
465              470              475              480

Thr Leu Thr Cys Thr Ala Glu Asn Gln Leu Glu Arg Thr Val Asn Ser
485              490              495

Leu Asn Val Ser Ala Ile Ser Ile Pro Glu His Asp Glu Ala Asp Glu
500              505              510

Ile Ser Asp Glu Asn Arg Glu Lys Val Asn Asp Gln Ala Lys Leu Ile
515              520              525

Val Gly Ile Val Val Gly Leu Leu Leu Ala Ala Leu Val Ala Gly Val
530              535              540

Val Tyr Trp Leu Tyr Met Lys Lys Ser Lys Thr Ala Ser Lys His Val
545              550              555              560

Asn Lys Asp Leu Gly Asn Met Glu Glu Asn Lys Lys Leu Glu Glu Asn
565              570              575

Asn His Lys Thr Glu Ala
580

<210> 335
<211> 709
<212> PRT
<213> Homo sapiens

<400> 335
Met Ala Glu Val Glu Asp Gln Ala Ala Arg Asp Met Lys Arg Leu Glu
5              10              15

Glu Lys Asp Lys Glu Arg Lys Asn Val Lys Gly Ile Arg Asp Asp Ile
20              25              30

Glu Glu Glu Asp Asp Gln Glu Ala Tyr Phe Arg Tyr Met Ala Glu Asn
35              40              45

Pro Thr Ala Gly Val Val Gln Glu Glu Glu Glu Asp Asn Leu Glu Tyr
50              55              60

Asp Ser Asp Gly Asn Pro Ile Ala Pro Thr Lys Lys Ile Ile Asp Pro
65              70              75              80

Leu Pro Pro Ile Asp His Ser Glu Ile Asp Tyr Pro Pro Phe Glu Lys
85              90              95

Asn Phe Tyr Asn Glu His Glu Glu Ile Thr Asn Leu Thr Pro Gln Gln
100             105             110

```

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Ile | Asp | Leu | Arg | His | Lys | Leu | Asn | Leu | Arg | Val | Ser | Gly | Ala | Ala |
| 115 | | | | | | 120 | | | | | | 125 | | | |
| Pro | Pro | Arg | Pro | Gly | Ser | Ser | Phe | Ala | His | Phe | Gly | Phe | Asp | Glu | Gln |
| 130 | | | | | | 135 | | | | | | 140 | | | |
| Leu | Met | His | Gln | Ile | Arg | Lys | Ser | Glu | Tyr | Thr | Gln | Pro | Thr | Pro | Ile |
| 145 | | | | | | 150 | | | | | | 155 | | 160 | |
| Gln | Cys | Gln | Gly | Val | Pro | Val | Ala | Leu | Ser | Gly | Arg | Asp | Met | Ile | Gly |
| | | | | 165 | | | | | | 170 | | | | 175 | |
| Ile | Ala | Lys | Thr | Gly | Ser | Gly | Lys | Thr | Ala | Ala | Phe | Ile | Trp | Pro | Met |
| | | 180 | | | | | | 185 | | | | | | 190 | |
| Leu | Ile | His | Ile | Met | Asp | Gln | Lys | Glu | Leu | Glu | Pro | Gly | Asp | Gly | Pro |
| | | 195 | | | | | | 200 | | | | 205 | | | |
| Ile | Ala | Val | Ile | Val | Cys | Pro | Thr | Arg | Glu | Leu | Cys | Gln | Gln | Ile | His |
| 210 | | | | | | 215 | | | | | | 220 | | | |
| Ala | Glu | Cys | Lys | Arg | Phe | Gly | Lys | Ala | Tyr | Asn | Leu | Arg | Ser | Val | Ala |
| 225 | | | | | | 230 | | | | | | 235 | | 240 | |
| Val | Tyr | Gly | Gly | Gly | Ser | Met | Trp | Glu | Gln | Ala | Lys | Ala | Leu | Gln | Glu |
| | | | | 245 | | | | | | 250 | | | | 255 | |
| Gly | Ala | Glu | Ile | Val | Val | Cys | Thr | Pro | Gly | Arg | Leu | Ile | Asp | His | Val |
| | | 260 | | | | | | 265 | | | | | | 270 | |
| Lys | Lys | Lys | Ala | Thr | Asn | Leu | Gln | Arg | Val | Ser | Tyr | Leu | Val | Phe | Asp |
| | | 275 | | | | | | 280 | | | | 285 | | | |
| Glu | Ala | Asp | Arg | Met | Phe | Asp | Met | Gly | Phe | Glu | Tyr | Gln | Val | Arg | Ser |
| 290 | | | | | | 295 | | | | | | 300 | | | |
| Ile | Ala | Ser | His | Val | Arg | Pro | Asp | Arg | Gln | Thr | Leu | Leu | Phe | Ser | Ala |
| 305 | | | | | | 310 | | | | | | 315 | | 320 | |
| Thr | Phe | Arg | Lys | Lys | Ile | Glu | Lys | Leu | Ala | Arg | Asp | Ile | Leu | Ile | Asp |
| | | | | 325 | | | | | | 330 | | | | 335 | |
| Pro | Ile | Arg | Val | Val | Gln | Gly | Asp | Ile | Gly | Glu | Ala | Asn | Glu | Asp | Val |
| | | 340 | | | | | | 345 | | | | | | 350 | |
| Thr | Gln | Ile | Val | Glu | Ile | Leu | His | Ser | Gly | Pro | Ser | Lys | Trp | Asn | Trp |
| | | 355 | | | | | | 360 | | | | 365 | | | |
| Leu | Thr | Arg | Arg | Leu | Val | Glu | Phe | Thr | Ser | Ser | Gly | Ser | Val | Leu | Leu |
| 370 | | | | | | 375 | | | | | | 380 | | | |
| Phe | Val | Thr | Lys | Lys | Ala | Asn | Ala | Glu | Glu | Leu | Ala | Asn | Asn | Leu | Lys |
| 385 | | | | | | 390 | | | | | | 395 | | 400 | |

Gln Glu Gly His Asn Leu Gly Leu Leu His Gly Asp Met Asp Gln Ser
 405 410 415
 Glu Arg Asn Lys Val Ile Ser Asp Phe Lys Lys Lys Asp Ile Pro Val
 420 425 430
 Leu Val Ala Thr Asp Val Ala Ala Arg Gly Leu Asp Ile Pro Ser Ile
 435 440 445
 Lys Thr Val Ile Asn Tyr Asp Val Ala Arg Asp Ile Asp Thr His Thr
 450 455 460
 His Arg Ile Gly Arg Thr Gly Arg Ala Gly Glu Lys Gly Val Ala Tyr
 465 470 475 480
 Thr Leu Leu Thr Pro Lys Asp Ser Asn Phe Ala Gly Asp Leu Val Arg
 485 490 495
 Asn Leu Glu Gly Ala Asn Gln His Val Ser Lys Glu Leu Leu Asp Leu
 500 505 510
 Ala Met Gln Asn Ala Trp Phe Arg Lys Ser Arg Phe Lys Gly Gly Lys
 515 520 525
 Gly Lys Lys Leu Asn Ile Gly Gly Gly Gly Leu Gly Tyr Arg Glu Arg
 530 535 540
 Pro Gly Leu Gly Ser Glu Asn Met Asp Arg Gly Asn Asn Asn Val Met
 545 550 555 560
 Ser Asn Tyr Glu Ala Tyr Lys Pro Ser Thr Gly Ala Met Gly Asp Arg
 565 570 575
 Leu Thr Ala Met Lys Ala Ala Phe Gln Ser Gln Tyr Lys Ser His Phe
 580 585 590
 Val Ala Ala Ser Leu Ser Asn Gln Lys Ala Gly Ser Ser Ala Ala Gly
 595 600 605
 Ala Ser Gly Trp Thr Ser Ala Gly Ser Leu Asn Ser Val Pro Thr Asn
 610 615 620
 Ser Ala Gln Gln Gly His Asn Ser Pro Asp Ser Pro Val Thr Ser Ala
 625 630 635 640
 Ala Lys Gly Ile Pro Gly Phe Gly Asn Thr Gly Asn Ile Ser Gly Ala
 645 650 655
 Pro Val Thr Tyr Pro Ser Ala Gly Ala Gln Gly Val Asn Asn Thr Ala
 660 665 670
 Ser Gly Asn Asn Ser Arg Glu Gly Thr Gly Gly Ser Asn Gly Lys Arg
 675 680 685

Ser Ser Tyr Glu Gln Asn Asp Asn Ser Leu Val Tyr Phe Ala Tyr Tyr
210 215 220

Met Ala Ala Ala Lys Ala Glu Met Gln Leu Met Ser Pro Leu Gln Ile
5 10 15

Ser Asp Pro Phe Gly Ser Phe Pro His Ser Pro Thr Met Asp Asn Tyr
20 25 30

Pro Lys Leu Glu Glu Met Met Leu Leu Ser Asn Gly Ala Pro Gln Phe
35 40 45

Leu Gly Ala Ala Gly Ala Pro Glu Gly Ser Gly Ser Asn Ser Ser Ser
50 55 60

Ser Ser Ser Gly Gly Gly Gly Gly Gly Gly Gly Gly Ser Asn Ser Ser
65 70 75 80

Ser Ser Ser Ser Thr Phe Asn Pro Gln Ala Asp Thr Gly Glu Gln Pro
85 90 95

Tyr Glu His Leu Thr Ala Glu Ser Phe Pro Asp Ile Ser Leu Asn Asn
100 105 110

Glu Lys Val Leu Val Glu Thr Ser Tyr Pro Ser Gln Thr Thr Arg Leu
115 120 125

Pro Pro Ile Thr Tyr Thr Gly Arg Phe Ser Leu Glu Pro Ala Pro Asn
130 135 140

Ser Gly Asn Thr Leu Trp Pro Glu Pro Leu Phe Ser Leu Val Ser Gly
145 150 155 160

Leu Val Ser Met Thr Asn Pro Pro Ala Ser Ser Ser Ser Ala Pro Ser
165 170 175

Pro Ala Ala Ser Ser Ala Ser Ala Ser Gln Ser Pro Pro Leu Ser Cys
180 185 190

Ala Val Pro Ser Asn Asp Ser Ser Pro Ile Tyr Ser Ala Ala Pro Thr
195 200 205

Phe Pro Thr Pro Asn Thr Asp Ile Phe Pro Glu Pro Gln Ser Gln Ala
210 215 220

Phe Pro Gly Ser Ala Gly Thr Ala Leu Gln Tyr Pro Pro Pro Ala Tyr
225 230 235 240

Pro Ala Ala Lys Gly Gly Phe Gln Val Pro Met Ile Pro Asp Tyr Leu
245 250 255

Phe Pro Gln Gln Gln Gly Asp Leu Gly Leu Gly Thr Pro Asp Gln Lys
260 265 270

Pro Phe Gln Gly Leu Glu Ser Arg Thr Gln Gln Pro Ser Leu Thr Pro

| | | |
|---|-----|---------|
| 275 | 280 | 285 |
| Leu Ser Thr Ile Lys Ala Phe Ala Thr Gln Ser Gly Ser Gln Asp Leu | | |
| 290 | 295 | 300 |
| Lys Ala Leu Asn Thr Ser Tyr Gln Ser Gln Leu Ile Lys Pro Ser Arg | | |
| 305 | 310 | 315 320 |
| Met Arg Lys Tyr Pro Asn Arg Pro Ser Lys Thr Pro Pro His Glu Arg | | |
| | 325 | 330 335 |
| Pro Tyr Ala Cys Pro Val Glu Ser Cys Asp Arg Arg Phe Ser Arg Ser | | |
| | 340 | 345 350 |
| Asp Glu Leu Thr Arg His Ile Arg Ile His Thr Gly Gln Lys Pro Phe | | |
| | 355 | 360 365 |
| Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp His Leu Thr | | |
| | 370 | 375 380 |
| Thr His Ile Arg Thr His Thr Gly Glu Lys Pro Phe Ala Cys Asp Ile | | |
| 385 | 390 | 395 400 |
| Cys Gly Arg Lys Phe Ala Arg Ser Asp Glu Arg Lys Arg His Thr Lys | | |
| | 405 | 410 415 |
| Ile His Leu Arg Gln Lys Asp Lys Lys Ala Asp Lys Ser Val Val Ala | | |
| | 420 | 425 430 |
| Ser Ser Ala Thr Ser Ser Leu Ser Ser Tyr Pro Ser Pro Val Ala Thr | | |
| | 435 | 440 445 |
| Ser Tyr Pro Ser Pro Val Thr Thr Ser Tyr Pro Ser Pro Ala Thr Thr | | |
| | 450 | 455 460 |
| Ser Tyr Pro Ser Pro Val Pro Thr Ser Phe Ser Ser Pro Gly Ser Ser | | |
| 465 | 470 | 475 480 |
| Thr Tyr Pro Ser Pro Val His Ser Gly Phe Pro Ser Pro Ser Val Ala | | |
| | 485 | 490 495 |
| Thr Thr Tyr Ser Ser Val Pro Pro Ala Phe Pro Ala Gln Val Ser Ser | | |
| | 500 | 505 510 |
| Phe Pro Ser Ser Ala Val Thr Asn Ser Phe Ser Ala Ser Thr Gly Leu | | |
| | 515 | 520 525 |
| Ser Asp Met Thr Ala Thr Phe Ser Pro Arg Thr Ile Glu Ile Cys | | |
| | 530 | 535 540 |

<210> 338

<211> 148

<212> PRT

<400> 338

Leu Phe Leu Thr Ile Pro Phe Ala Phe Phe Leu Pro Glu Leu Ile Phe
20 25 30

Gly Phe Leu Val Trp Thr Met Val Ala Ala Thr His Ile Val Tyr Pro
35 40 45

Leu Leu Gln Gly Trp Val Met Tyr Val Ser Leu Thr Ser Phe Leu Ile
50 55 60

Ser Leu Met Phe Leu Leu Ser Tyr Leu Phe Gly Phe Tyr Lys Arg Phe
65 70 75 80

Glu Ser Trp Arg Val Leu Asp Ser Leu Tyr His Gly Thr Thr Gly Ile
85 90 95

Leu Tyr Met Ser Ala Ala Val Leu Gln Val His Ala Thr Ile Val Ser
100 105 110

Glu Lys Leu Leu Asp Pro Arg Ile Tyr Tyr Ile Asn Ser Ala Ala Ser
115 120 125

Phe Phe Ala Phe Ile Ala Thr Leu Leu Tyr Ile Leu His Ala Phe Ser
130 135 140

Ile Tyr Tyr His
145

<210> 339

<211> 196

<212> PRT

<213> Homo sapiens

<400> 339

Met Pro Gly Met Phe Phe Ser Ala Asn Pro Lys Glu Leu Lys Gly Thr

5 10 15

Thr His Ser Leu Leu Asp Asp Lys Met Gln Lys Arg Arg Pro Lys Thr
20 25 30

Phe Gly Met Asp Met Lys Ala Tyr Leu Arg Ser Met Ile Pro His Leu
35 40 45

Glu Ser Gly Met Lys Ser Ser Lys Ser Lys Asp Val Leu Ser Ala Ala
50 55 60

Glu Val Met Gln Trp Ser Gln Ser Leu Glu Lys Leu Leu Ala Asn Gln
65 70 75 80

[illegible]

```
<210> 342
<211> 472
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1) ... (472)
<223> n = A,T,C or G
```

```
<210> 343
<211> 139
<212> DNA
<213> Homo sapien
```

```
<210> 344
<211> 235
<212> DNA
<213> Homo sapien
```

```
<210> 345
<211> 458
<212> DNA
<213> Homo sapien
```

| | | | | | | |
|-------------|-------------|------------|-------------|------------|------------|-----|
| <400> 345 | | | | | | |
| ctgtaagggtg | ctatttcagtc | ctgtgaccct | tatttttgaa | tgctcttcat | tactgttget | 60 |
| ctgtttttgtg | acttcctggg | aaaccgccta | ctttgggtgtg | gtgtcacctt | gagctgtgca | 120 |
| cataggacac | cagtttttgac | ttaacctaac | aggcagtttt | tatctctagc | tttttcaagc | 180 |

```
<210> 346
<211> 525
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(525)
<223> n = A,T,C or G
```

```
<210> 347
<211> 423
<212> DNA
<213> Homo sapien
```

```
<210> 348
<211> 513
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|------------|------------|------------|-------------|--|-----|
| <400> | 348 | | | | | | |
| cctctaggcc | tgatgctctc | agaggcaata | gaagaaaagt | aaaaggaagg | tctcacttca | | 60 |
| cagacaatga | aaccctccta | accctcttcc | ccactaccca | caactcccta | cactgccaat | | 120 |
| ctaaataaaa | agaggacaat | gcatgagtg | gagatacaca | tacacacaca | cacatacaca | | 180 |
| cacacacacg | cacagcttcc | tttcagccaa | agaactgcaa | aatccttccc | cgggaaggagg | | 240 |
| acaactggca | acaccaatca | aggcttgg | gtctaagg | atggctggaa | tcattgtgaga | | 300 |
| ctggtaaaaa | tccagggaga | aaatgtttca | ccttcagctc | attcccaagt | ctctatgaag | | 360 |

```

cccgccccac ttccacatag gggaaactgtg gctctggggg cagcctctgc agctactcag      420
aataggtggg aggaggggct ggctttgagg ctgccttagc catgaggctc tttgcctagg      480
aatagctgga gatgggagct gcagggggct cag                                     513

```

```

<210> 349
<211> 231
<212> DNA
<213> Homo sapien

```

```

<400> 349
ccttatttct cttgtccttt cgtacagggg ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta      120
atagcggtcg caccatcggt atgtcctgat ccaacatcga ggtcgtaaac cctattgttg      180
atatggactc tagagtagga ttgcgctggt atccctaggg taacttgttc c                231

```

```

<210> 350
<211> 341
<212> DNA
<213> Homo sapien

```

```

<400> 350
ctgcccaagg gcgttcgtaa cggaatgcc gaagcgtggg aaaaaggagg cgggtggcgga      60
agacggggat gagctcagga cagagccaga ggccaagaag agtaagacgg ccgcaaagaa      120
aatgacaaa gaggcagcag gagagggccc agccctgtat gaggaccccc cagatcagaa      180
aacctcacc agtggcaaac ctgccacacc caagatctgc tcttggaatg tggatgggct      240
tcgagcctgg attaagaaga aaggattaga ttgggtaaag gaagaagccc cagatatata      300
gtgccttcaa gagaccaaat gttcagagaa caaactacca g                    341

```

```

<210> 351
<211> 256
<212> DNA
<213> Homo sapien

```

```

<400> 351
ggcgttgggg acggtttagt gacgtggctc tttattcgtg agttttccat ttacctccgc      60
tgaacctaga gcttcagacg ccctatggcg tccgcctcga cccaaccggc ggccttgagc      120
gctgagcaag caaagggtgt cctcgcggag gtgatccagg cgttctccgc cccggagaat      180
gcagtgcgca tggacgaggc tcgggataac gcctgcaacg acatgggtaa gatgctgcaa      240
ttcgtgctgc ccgtgg                                     256

```

```

<210> 352
<211> 368
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(368)
<223> n = A,T,C or G

```

```

<400> 352
cctttcttgt aagtgaagaa naaggaatgc agcaaagaag agttcgacat tggagtcctt      60
agttccatca ggatcccatt cgcagccttt agcatcatgt agaagcaaac tgcacctatg      120
gctgagatag gtgcaatgac ctacaagatt ttgtgttttc tagctgtcca ggaaaagcca      180

```


tcttcagtct tgctgacagt caaagagcaa gtgaaacccat ttccagccta aactacataa 240
aagcagccga accaatgatt aaagacctct aaggctccat aatcatcatt aaatatgcc 300
aaactcattg tgacttttta ttttatatac aggattaaaa tcaacattaa atcatcttat 360
ttacatgg 368

<210> 353
<211> 368
<212> DNA
<213> Homo sapien

<400> 353
ctgaggggtg gcagtaagca atgaggatgg gctataaagc tgttaactgg ctaagggcca 60
tccttgggca ggcatttcag acacatctgt agagagggca gtagcatctc cgataggcca 120
gctctgaagg aagcttaatg ctttaatacag tcacactgca taaattagct tagaatgctc 180
tcttgggtaa aaaatattaa tagtgtatat gcacttgaag agcaaaattc ctcaagaaaa 240
aaagttaaag agcaaggagt ttccatcagt cccgggtctt gtgaggatta ccacaacaaa 300
cacttaaaag gatacaacag gtacttatta aatgctgcct tgccttttac ctcttccttt 360
tttttttt 368

<210> 354
<211> 380
<212> DNA
<213> Homo sapien

<400> 354
ccatggcttc tcacccagac agtctttctg ggcaacttgg ggaagccctt gttctgctca 60
agtctcacc catggaagag gtgggggaag ggggccttgg tttttcagga agacagggtg 120
gagagcacga gtcactacaa agcagtaaaa gtgaatggtg tctccagggg ctgggtccag 180
aacaccacgg agagccccag ccataaagggt gtgttccgcc tctggcctgc aggaatctct 240
ttgaatctct ttgattggtg gctccaagag caatgggaag tcaacagcca ggaggctgga 300
ctgggttccc tgggaccccg aggtcccaga gctgctgggc agtggttgct ggcaaagaag 360
aaagggtcaa gagggtcagg 380

<210> 355
<211> 347
<212> DNA
<213> Homo sapien

<400> 355
ccagtggagg ggtgggggta tcgatcccg cgggggctgg cttggttgct ggtgcctga 60
gcccttctct gccgcctgg gtgttgctt cactgatgga ggtagcgct cagccagatg 120
tcaccagact tcttcgggga cctgacgatg tccaccagcg cggtgaggaa gggttcaact 180
tcgtagctga ggccgtgctt ggcacacagc gacttgacca gcggggccac ccggctgtag 240
ttgtgtctcg gcactctggg gaagaggtgg tgctcgatct ggaagttgag gtgcccgcgtg 300
aaccagttgg tgaaaagtga gggctccacg ttgcaggtgg ctgccag 347

<210> 356
<211> 157
<212> DNA
<213> Homo sapien

<400> 356
cctggagctg ctgaagactg ctattgggaa agctggctac actgataagg tggatcatcg 60
catggacgta gcggcctccg agttcttcag gtctgggaag tatgacctgg acttcaagtc 120

157

<400> 357

```
<210> 358
<211> 555
<212> DNA
<213> Homo sapien
```

<400> 358

```
<210> 359
<211> 549
<212> DNA
<213> Homo sapien
```

<400> 359

```
<210> 360
<211> 289
<212> DNA
<213> Homo sapien
```

<400> 360
 tttaaatttt actagtgtta cttaattgtat atttctaaaaa gagaatgcag taactaatgc 60
 cctaaatggt tgatctctgt ttgtcattac tttttcaaaa ttattttttt ctgtaaagta 120
 taatatataa aacttcttgc ttaaattgaa tttctatatt agtggttaat tgcagtttat 180
 taaagggatc attatcagta atttcatagc aactgttcta gtgttttgtg tttttaaaac 240
 agaattagga atttgagata tctgattata tttttcatat gaatcacag 289

<210> 361
 <211> 311
 <212> DNA
 <213> Homo sapien

<400> 361
 ctgttcagta tggcaaaggg cagacttact ctttcatcca ctctgctgcc ttgatgaggt 60
 gaacacactg gaataagatg gagggcagga tacctgccaa agcctgagga atgagatgat 120
 ctgaaacaat tgggcaaagg ctggacattt caaaaagctg acttccaact gcagtttatg 180
 ggtatagaat ttgatgtctt cctcaagtcc tgactgctct ttctgaggca gccaggctag 240
 gccaagaaat gagctgctcc agcttctcca gagcacagca gcctcccagg gcctgtcagc 300
 atctgcagca g 311

<210> 362
 <211> 496
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(496)
 <223> n = A,T,C or G

<400> 362
 ccagtttcta aaanaatgca catttaaaga gaagcatcta ccacggcttt aaaacaaaac 60
 aactctgaga tgaacaatat gtgttatact cagagattaa caatctcaat catacatact 120
 gattctttca gacatttaac aaccactaca tttttttgca ttaatgaagt ttgactatat 180
 gtgtaaaggg actaaatatt tttgcaacag cctgttcttt gttcattctt ttctggatag 240
 cgtgtcctct gtattgcggt agatttatac attctgttgc ctaaatatgt gtgtaaaatg 300
 agctgataaa ctggagtact acttaaaaaa aagtctgtga tttataagat gcatatgctt 360
 tctatgtgaa tataagcttg tgcacaatgt ttaaaagaaa aacaatgaat tagaagagat 420
 cccccgtccc ccagtctgac atatttcata cagaatgttt aaaagaaaaa ctctgctagt 480
 cttggcaaac atttgg 496

<210> 363
 <211> 673
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(673)
 <223> n = A,T,C or G

<400> 363
 ccaagaggga gataanacaa acttctcaaa caaaaagaaa agaaaaacga atgattcatc 60
 tgctttaatc agtgtgatta atgcagcacc cattgccccg ggaaccgttt ctgctgtact 120

```

atctggatac taaaatgtta cggaagtagc tctttgttct ccctcactct gcccttagtt 180
aatagaaatt cagactcgcc aagtaaggct ttgtgcatag tgtcttcatt tcgcgtatag 240
ttgagcgcgt tcttagcagt tggcttcatt gacagctcat tagtgttttg acttttctta 300
cccagcgtta attgaattct tgcttttaga caacttcctt tttgtagtgg tgaaccttgc 360
ccttttagtac agttcaagtg aatctggata attgttcatt tttgctttag cttagatacc 420
atgtagtggg ctgtggctac aggaagctgg ttctgtctgc ttccacagtc tgcttaaaaa 480
actgtctgac ttcgtgaata tagagaccaa gtttaccact tctgatgaag agaccaatta 540
agattcattc ctcatctgtt ttctttccag tgggagaaga gtcccatga aataagatga 600
aactgattcc atgcactagt acatgtaggc ttctcccttg cgcaaagctt aacaatttgt 660
aggaaacttt ggg 673

```

```

<210> 364
<211> 495
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(495)
<223> n = A,T,C or G

```

```

<400> 364
ccaaatgttt gcncaagact agcagagttt ttcttttaaa cattctgtat gaaatatgtc 60
agactggggg acgggggatc tcttctaatt cattgttttt cttttaaaca ttgtgcacaa 120
gcttatattc acatagaaag catatacatc ttataaatca cagacttttt ttttaagtagt 180
actccagttt atcagctcat ttacacaca tatttaggca acagaatgta taaatctacc 240
gcaatacaga ggacacacta tccagaaaag aatgaacaaa gaacaggctg ttgcaaaaat 300
atttagtccc ttacacata tagtcaaact tcattaatgc aaaaaatgta gtggttatta 360
aatgtctgaa agaatcagta tgtatgattg agattgttaa tctctgagta taacacatat 420
tgttcatctc agagttgttt tgttttaaaag ccgtggtaga tgcttctctt taaatgtgca 480
tttttttagaa actgg 495

```

```

<210> 365
<211> 291
<212> DNA
<213> Homo sapien

```

```

<400> 365
aactgacaag cccttgccgc tgctctcca ggatgtctac aaaattggtg gtattggtac 60
tgttcctgtt ggcccagtg gagactggtg ttctcaaacc cggtatggtg gtcacctttg 120
ctccagtcac cgttacaacg gaagtaaaat ctgtcgaaat gcaccatgaa gctttgagtg 180
aagctcttcc tggggacaat gtgggcttca atgtcaagaa tgtgtctgtc aaggatgttc 240
gtcgtggcaa cgttgctggt gacagcaaaa atgacccacc aatggaagca g 291

```

```

<210> 366
<211> 277
<212> DNA
<213> Homo sapien

```

```

<400> 366
ctggatgggt cctcagaagg tgcattctgc ttctgcaggg gcttgaaaca ccaaggcact 60
ccagggatcc tggagtcaaa gcagcagccc cggttggtgc actccttggg ggtgacatgg 120
gggtagcccc cagtcacccc tgtccttggc tggcacggca cactggtttg cagacaggcc 180
cacgtactcc tcagcagagc tggaggacaa gcaaggccag gaccagcccc agcatgcaga 240

```

gcgctctggc agccatgacc accgtgggct ccgggac

277

<210> 367
 <211> 311
 <212> DNA
 <213> Homo sapien

<400> 367
 ccagagctgc ggggcctcag tacacggagc tgttccggat gccacagcac agcaccatgc 60
 tcaggatcat ctogaagatc atgatacacg cgaccacgat ggcagcaatg ccgatgaggt 120
 acagcttccc ggagaagagg tcatcgatct tctgggtggca gtcctccttg aagaggttgc 180
 tgatgatgtt gctgcccagag ggacacaaat tgttcttgag cactgaggtg gtcaaagcag 240
 tcagtgtgct ggagccacag cagtcaagcg tctcgtggaa ggtcttcacc acagccttgg 300
 cgttgttggc g 311

<210> 368
 <211> 384
 <212> DNA
 <213> Homo sapien

<400> 368
 ccaaaggggt ctctagctgc tgcctctgctg ctccctgctca tggatgagtt tggcgatggg 60
 gccggtgatg ccgcctatca aggtccagta ctcatcgaag ctgatgcgcc catcaggatt 120
 ggcattccagg ttctggatga gcttatccgc agccttccgg ttccctgtgt ccgacagcat 180
 gtggttcagc tctttctgga gcatctcgcg gaagctgctc ttgctgatct tgttcttgac 240
 caggctgtac ctagacacat attttagtaa gttttccacc aggacaatga ctgccttctc 300
 cagctccgtg tagcaagtct gacatctccc tgcttgcctt gctggcgggg cctaaggcgg 360
 gggccaagcc cagttacagc ccag 384

<210> 369
 <211> 216
 <212> DNA
 <213> Homo sapien

<400> 369
 ccaagtgcc a ggtggctttc agcagcttcc tacgatcagc cgaagaaagc agaagctctg 60
 gaggtgcc a tcgagaacct caatgaagcc aagaactatt ttgcaaagg t gactgcaaa 120
 gagcgcatca gggacgtcgt ttacttccag gccagactct accataccct ggggaagacc 180
 caggagagga accggtgtgc gatgctcttc cggcag 216

<210> 370
 <211> 561
 <212> DNA
 <213> Homo sapien

<400> 370
 ctggctcctt cttttgtggt cgtttggggg atgggctggt ttggggttta ggtgcagaga 60
 atggtttggg gccactgcgt actggaccac tctgagcctt cagggcaggg ttcttgtgag 120
 tcttcatgtc atcagataca tgtttcaggg catgtgtaat gctctcccc tgattaatct 180
 ggcgaacag tctgagcgg gaagcagact catctgagcc tgaactggta gagactgggg 240
 gaggaggggg gcttgggtgga gggggaggag gacctgatcc ggcagagggg ccagatggca 300
 gtccgctcag ttcttttggc acaggcccc ttttgcctca ggccagtcgg gtggtatgga 360
 actccttaat gtaagcctgc agctctgtcc atatacttaa ataagctttg acccagtcta 420
 catgcttctt atccacatct ttgtactctt tgaggactcg gtttgtataa aacatggcgg 480

```
<210> 371
<211> 518
<212> DNA
<213> Homo sapien
```

```
<210> 372
<211> 335
<212> DNA
<213> Homo sapien
```

```
<210> 373
<211> 467
<212> DNA
<213> Homo sapien
```

```
<210> 374
<211> 284
<212> DNA
<213> Homo sapien
```

<400> 374
ttttccgtataa agcgtgtaac aaggggtgtaa atatttataa ttttttatac ctgtttgtgag 60

```
<210> 375
<211> 307
<212> DNA
<213> Homo sapien
```

```
<210> 376
<211> 650
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(650)
<223> n = A,T,C or G
```

```
<210> 377
<211> 306
<212> DNA
<213> Homo sapien
```

```
<220>  
<221> misc_feature  
<222> (1)...(306)  
<223> n = A,T,C or G
```

<400> 377
tctagatgca tgctcgagcg gccgccagtg tgatgganat ctgcagaatt cgcccttoga 60
gcggccgccc gggcaggttc ggggtgctgcc ttcacctgcc aggcccttcc ccgctagett 120

```
<210> 378
<211> 199
<212> DNA
<213> Homo sapien
```

<400> 378

```
<210> 379
<211> 216
<212> DNA
<213> Homo sapien
```

<400> 379

```
<210> 380
<211> 555
<212> DNA
<213> Homo sapien
```

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccatgggcct | tcccttccac | taaaaggaat | tccgaacagc | aaaaagaagg | tcttgagata | 60 |
| gtgaaaatgg | tgatgatatc | tttagaaggt | gaagatgggt | tggatgaaat | ttattcattc | 120 |
| agtgagagtc | tgagaaaact | gtgcgtcttc | aagaaaattg | agaggcattc | cattcactgg | 180 |
| ccctgcgcac | tgaccattgg | ctccaatttg | tctataagga | ttgcagccta | taaatcgatt | 240 |
| ctacaggaga | gagttaaaaa | gacttgga | gttggtgatg | caaaaaccct | aaaaaaagaa | 300 |
| gatatacaaa | aagaaacagt | ttattgctta | aatgatgatg | atgaaactga | agttttaaaa | 360 |
| gaggatatta | ttcaagggtt | ccgctatgga | agtgatatag | ttcctttctc | taaagtggat | 420 |
| gaggaacaaa | tgaaatataa | atcggagggg | aagtgccttc | ctgttttggg | attttgtaaa | 480 |
| tcttctcagg | gtcagagaag | attcttcatg | ggaaatcaag | ttctaaaggc | tttgcccaa | 540 |
| gagatgatga | ggcag | | | | | 555 |

<210> 381
 <211> 406
 <212> DNA
 <213> Homo sapien

```

<400> 381
ctgcaccagg tgggcctcta ggtcccatta agcccattgg tccagggcca agtccaactc      60
cttttccatc atactgagca gcaaagttcc caccgagacc agggggggcca ggaggaccag      120
gtggaccagg agggcctgtg ggaccatctt caccatctct gcctgggggg cctgggtggac      180
ccctttctcc acgtggtcct ctatctccgg ctgggccctt tcttacagtt tcctcttgta      240
aagattggca tgttgctagg cataaggtta ctgcaagcag caacaaagtc cgcgtatcca      300
caaagctgag catgtctagc acttagacat gcgactcct tgtgtcgcag agcccctggg      360
tcaccggcgg aggtatcacc tggcgggcgc gggcatgcag tcgtgg                        406

```

<210> 382
 <211> 528
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(528)
 <223> n = A,T,C or G

```

<400> 382
ctgagcagtt tgtgggtntn tcttcccgcg agtttcagga agtattcaca aaagaaaaat      60
acattttttc ccccgagggg ggggcaagga cagtggagag agtgctagga aatgagtccc      120
ctgggaaagg ggaccgggccc gtgatgttaa atatctccgg ctccaagtg actggatttg      180
cctaggacct tcagaccaac agacttcaga cctcagacc tgccccgggg ccagggtggag      240
aaagtgaggg cgtacaagg aagtgaatt ctgagttggt ggggctaagc ctgacccctt      300
ctccatgctc cccgccccaa cccactctgg cctcagtaga tttttttttc agttgtggtt      360
gttgcccagg ctggagtgca gtagcgccat cttggctcac tgcacctcca ccttcggggc      420
tcaagcgatt ctccagctc agcctcctga gtagctagga ctgcaggtgc tccaccacgc      480
ccggctaatt tttgtatttt tagtagagat ggggtttccc catgttgg                        528

```

<210> 383
 <211> 335
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(335)
 <223> n = A,T,C or G

```

<400> 383
ccatnttgag tctactcctg cgtcttgtgc cctagcaccc cgagaaccgt cagtttgagc      60
cagatggaag ctgagctgaa cacattacga tggatgatgg aaacataaga ctatcaagaa      120
atccaagtgg taatgggcga agttttattca gcatccggca atggacttat cgtagttggg      180
gaaacgggtg ttccgaataa tatcctggaa gttatcagga cacctatttt aaatataggc      240
ctgaattttg taaagtaata tttaagggtg tccgtgataa ttaaataaaa tgcttaattc      300
atgtggcgaa aaaaaaaaaa naaaaaaaaa aaaaaa                        335

```

<210> 384

ggtgg

365

<210> 392
 <211> 302
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(302)
 <223> n = A,T,C or G

<400> 392
 ccaagagcta caatgagcag cgcatacanga cagaacgtgc aggtttttga gttccagttg 60
 actgcagagg acatgaaagc catagatggc ctagacagaa atctccacta ttttaacagt 120
 gatagttttg ctagccaccc taattatcca tattcagatg aatattaaca tggagagctt 180
 tgcctgatgt ctaccagaag ccctgtgtgt ggatggtgac gcagaggacg tctctatgcc 240
 ggtgactgga catatcacct ctacttaaat ccgctoctgtt tagcgacttc agtcaactac 300
 ag 302

<210> 393
 <211> 213
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(213)
 <223> n = A,T,C or G

<400> 393
 ccaataatca agnacaaana ctggatttga ggatggatca gttctgaaac agtttctttc 60
 tgaaacagag aaaatgtccc ctgaagacag agcaaaatgc tttggaaaga atgaggccat 120
 acaggcagcc catgatgccg tggcacagga aggccaatgt cgggtagatg acaagggtgaa 180
 tttccatttt attctgttta acaacgtgga tgg 213

<210> 394
 <211> 334
 <212> DNA
 <213> Homo sapien

<400> 394
 cctacccata atccagagag gcttgcccag aggaggacta cgtggggggac gtgccaccag 60
 aaccctactt gggggcgagg tgtcactccg aggtcaaaac ctgctccgag gtggacgagc 120
 cgtagctccc cgaatgggct taagaagagg tgggtgtcga ggtcgtggag gtcctgggag 180
 agggggccta gggcgtggag ctatgggtcg tggcggaaac ggtggttagag gtcgggggtat 240
 gataggctcg ggaagagggg gctttggagg ccgaggccga ggccgtggac gagggagagg 300
 tgcccttgct cgccctgtat tgaccaagga gcag 334

<210> 395
 <211> 174
 <212> DNA
 <213> Homo sapien

<210> 399
 <211> 329
 <212> DNA
 <213> Homo sapien

<400> 399
 ccaacctcag gcaacgggtg gagcagtttg ccagggcctt ccccatgcct ggttttgatg 60
 agcattgaag gcacctggga aatgaggccc acagactcaa agttactctc cttcccccta 120
 cctggggcag tgaaatagaa agcctttcta ttttttggtg cgggagggaa gacctctcac 180
 ttagggcaag agccaggtat agtctccctt cccagaatth gtaactgaga agatcttttc 240
 tttttccttt tttcggtaac aagacttaga aggagggccc aggcactttc tgtttgaacc 300
 cctgtcatga tcacagtgtc agagacgcg 329

<210> 400
 <211> 451
 <212> DNA
 <213> Homo sapien

<400> 400
 ctggcttcac tgctcaggtg attatcctga accatccagg ccaaataagc gccggctatg 60
 cccctgtatt ggattgccac acggctcaca ttgcatgcaa gtttgctgag ctgaaggaaa 120
 agattgateg ccgttctggt aaaaagctgg aagatggccc taaattcttg aagtctggtg 180
 atgctgccat tgttgatatg gttcctggca agcccatgtg tgttgagagc ttctcagact 240
 atccaccttt gggtcgcttt gctgttcgtg atatgagaca gacagttgag gtgggtgtca 300
 tcaaagcagt ggacaagaag ctgctggagc tggcaaggct accaagtctg cccagaaagc 360
 tcagaagcta aatgaatatt atccctaata cctgccaccc cactcttaat cagtgggtgga 420
 agaacggctc agaactgttt gtttcaattg g 451

<210> 401
 <211> 180
 <212> DNA
 <213> Homo sapien

<400> 401
 ccaggaagca ggccagggga ttggcagcac tgcccagcac cacagccagg ttgtaggcca 60
 gacgcccgtg gggtaagcag gaaaagctct gcacggcagg cagcacgcca ttggtcagcg 120
 cgttgggtggc ggccaacagg cccagcaggc aggcaactgcg ggctgataga agctgatagg 180

<210> 402
 <211> 385
 <212> DNA
 <213> Homo sapien

<400> 402
 ccagggccacc tgtgcggggc tcctcgatgt ggaaggttcg ggtgaggaga ttgtagaagg 60
 agccgtagca cacggccacc acagtgcacg tgaggcagat cacgttgtag ggcatgctga 120
 agtccggtgt cggcaggttc accagcagcg gctccgtgta gagccgcaca aagtagttag 180
 agccatcaga gactgggaac aggctgttga agaggggact ctcttcccag tccactggct 240
 tggctgctac catgctgggc acaagggcgc tgaggacaga tgggctgaca tagaagccat 300
 ggtaggtatc tggcgtgtac tcggtccact tcagcagcgc ccgctcaaac tggatggaaa 360
 ccttggtgac tgagttggcc ggcag 385

<210> 403
 <211> 440

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(440)
<223> n = A,T,C or G

<400> 403
ctgtttaacc agnaacccgg ggggtcacc cccacagaat gtacatgaaa cactagagga 60
ctgcatgttt ttccctgaga gaagcgtaag acaaacagaa gtcaaaaagt agtcactggg 120
agcgccatcc ttctaagcaa atcctccctt tcccttttgg aggatttgcc cgaactacgt 180
agccagtcag cacttagacc acctgcctcc tccccccct ataaaccac cactccctc 240
ctcctttccc aaaccacttg ggggtgccta agccctcact gcccgaagcc caaaatatca 300
gctaagatcc ttgtcagtat ttccacagtc atacctaatg aattgggaag tggggcccct 360
aaaaaccaat tcacatctat gcacttggtt ccactggatt tggcagacag gcttttttag 420
ttaccgtaac cagatcttaa 440

<210> 404
<211> 239
<212> DNA
<213> Homo sapien

<400> 404
cctacgaaaa actcccggcc ggtgaagaga acgtcagtgc catccagcgt cgcgttctcg 60
tctcctatct ccacaattcg gagccccagg tcttgacagg ctttgoggac tccatcgacc 120
tctggcctac gagcggggct ccagggccgc gtgattaggg ccgtgtcccc ttggatcacg 180
gccgtgtcgc caagcagcgg tcccagcggc aatgactcct cagggtggcag ttctagcag 239

<210> 405
<211> 261
<212> DNA
<213> Homo sapien

<400> 405
ctggagaggc agcccttcac cggatgccc gctccgtgcc cctgcggggc ccagcacagt 60
ttaccttctc cccccacggc ggtcccatct actctgtgag ctgttcccc ttccacagga 120
atctcttctt gagecgtggg actgacgggc atgtccacct gtactccatg ctgcaggccc 180
ctcccttgac ttgcgtgcag ctctccctca agtatctgtt tgcgtgtgcg tgggtccccag 240
tgcgccctt ggtttttgca g 261

<210> 406
<211> 641
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(641)
<223> n = A,T,C or G

<400> 406
ctgctcccg gcntgggtggc agcaagtaga catcgggcct gtgcaggggc acccccttgg 60
gccgggagat ggtctgcttc agtggcgagg gcaggtctgt gtgggtcacg gtgcacgtga 120

006230" E3F360


```

ccatcagnga gaaaggtggt tgtcagttgt ttcacaaacc agattgagga ggacaaactg      60
ctctgccaat ttctggattt ctttattttt agcaaacact ttctttaag cttgactgtg      120
tgggcactca tccaagtgat gaataatcat caagggtttg ttgcttgtct tggatttata      180
tagagctttt tcatatgtct gagtccagat gagttggtca cccaacctc tggag          235

```

```

<210> 411
<211> 294
<212> DNA
<213> Homo sapien

```

```

<400> 411
aattaaggga agatgaagat gataaaacag ttttggatct tgctgtggtt ttgtttgaaa      60
cagcaacgct tcggtcaggg tatcttttac cagacactaa agcatatgga gatagaatag      120
aaagaatgct tcgcctcagt ttgaacattg accctgatgc aaaggtggaa gaagagcctg      180
aagaagaacc tgaagagaca gcagaagaca caacagaaga cacagagcaa gacgaagatg      240
aagaaatgga tgtgggaaca gatgaagaag aagaaacagc aaaggaatct acag          294

```

```

<210> 412
<211> 433
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(433)
<223> n = A,T,C or G

```

```

<400> 412
cctgagaagc cagaggcagg tggagagggg gtggaaagtg agcagcgggc tgggctggag      60
ccgcacacgc tctcctccca tgtaaataag cacctttaga aaaattcaca agtccccatc      120
cacaaaaaaa aaaanaanaa aaatttcagg gantaaaaat anactttgaa caaaaaggaa      180
catttgntgg cctggggggg catctnantt tntntagcnc cagngattcc ctccccnccc      240
cacccatcac atanatgtaa cacctttggt ntaaaatggg gagccgtttc caccntgccc      300
ccntccccgc cccagggcag ttgccccggn gacacntcaa gacaggancg aggtagtntt      360
tcancancac agttncacaa ggaacagaac agtntctccc gccagccct gcggcacaag      420
ggattgacac gcn          433

```

```

<210> 413
<211> 494
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(494)
<223> n = A,T,C or G

```

```

<400> 413
ccttattttct cttgtcnctt cgtacagggg ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta      120
atagcggctg caccatcggg atgtcctgat ccaacatcga ggtcgtaaac cctattgttg      180
atatggactc tagaatagga ttgcgctggt atccctaggg taacttgttc cgttggtcaa      240
gttattggat caattgagta tagtagttcg ctttgactgg tgaagtctta gcatgtactg      300
ctcggagggtt gggttctgct ccgaggtcgc cccaaccgaa atttttaatg cagggttggt      360

```

```

agtttaggac ctgtgggttt gttagggtact gtttgcatta ataaattaaa gtcctatagg 420
gtcttctcgt cttgctgtgt tatgcccgcc tcttcacggg cagggtcaatt tcaactggta 480
aaagtaagag acag 494

```

```

<210> 414
<211> 294
<212> DNA
<213> Homo sapien

```

```

<400> 414
ctgggcggat agcaccgggc atatttttga atggatgagg tctggcacc tgagcagtcc 60
agcgaggact tggctcttagt tgagcaattt ggctaggagg atagtatgca gcacggttct 120
gagtctgtgg gatagctgcc atgaagtaac ctgaaggagg tgctggctgg taggggttga 180
ttacagggtt gggaacagct cgtacacctg ccattctctg catatactgg ttagtgaggt 240
gagcctggcg ctcttctttg cgtgagcta aagctacata caatggcctt gtgg 294

```

```

<210> 415
<211> 421
<212> DNA
<213> Homo sapien

```

```

<400> 415
ccttgccccct gccctccac gaatgggttaa tatatatgta gatatatatt ttagcagtga 60
cattcccaga gagccccaga gctctcaagc tcctttctgt cagggtgggg ggttcagcct 120
gtcctgtcac ctctgagggt cctgctggca tcctctcccc catgcttact aatacattcc 180
cttccccata gccatcaaaa ctggaccaac tggcctcttc ctttccccctg ggacccaaat 240
ttaggggcct cagtcctca ccgccatgcc ctggcctatt ctgtctctcc ttcttcccc 300
tggcctgttc tgtctctgag ctctgtgtcc tccgttcatt ccatggctgg gagtcaactga 360
tgctgcctct gccttctgat gctggactgg ccttgcttct acaagtatgc ttctcccaca 420
g 421

```

```

<210> 416
<211> 342
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (342)
<223> n = A,T,C or G

```

```

<400> 416
ccactttctt tcccacnctg gaaggcggca tctatgactt cattggggag ttcatgaagg 60
ccagcgtgga tgtggcagac ctgataggtc taaaccttgt catgtcccgg aatgccggca 120
agggagagta caagatcatg gttgctgccc tgggctgggc cactgctgag cttattatgt 180
cccgtgcac tcccctatgg gtcggagccc ggggcattga gtttgactgg aagtacatcc 240
agatgagcat agactccaac atcagtctgg tccattacat cgtcgcgtct gctcagggtc 300
ggatgataac acgctatgat ctgtaccaca ctttccggcc gg 342

```

```

<210> 417
<211> 389
<212> DNA
<213> Homo sapien

```

<400> 417

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tattaattag | gttcttaaga | catttagaac | accaatttgt | gaggataaat | tccattcgtc | 60 |
| agagcaaaca | cagatcgcag | gtagccctgg | agctgaggaa | tagctttgat | ttttggtaaa | 120 |
| atthgtgagt | ccacagcttt | ctgatcaatc | ttgcgctgct | ccgtaatctc | atatttctct | 180 |
| ttttctgtgt | cgaagatctc | accttcctgg | tgtctgggct | tccgcagctt | cttcttcttg | 240 |
| aagtaagcat | cagtaagatg | ttttgggatt | tttacattgc | tgatatcgat | tttggttgaa | 300 |
| gtggcaatga | caaatttctg | gtgtgttctt | cgtagaggaa | ctcgattgag | gaccagaggt | 360 |
| ccagtcacaa | gtaataagcc | actagccag | | | | 389 |

<210> 418

<211> 343

<212> DNA

<213> Homo sapien

<400> 418

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| gtgggagggg | gccaggttgg | gatggagggg | gtttacagga | agcagacagg | gccaacgtcg | 60 |
| aagccgaatt | cctggctctg | ggcaccaacg | tccaaggggg | ccacatcgat | gatgggcagg | 120 |
| cgggaggtct | tggtggtttt | gtattcaatc | actgtcttgc | cccaggctcc | gggtgtgactc | 180 |
| gtgcagccat | cgacagtgc | gctgtagggt | aagcggctgt | tgccctcggc | gcggatctcg | 240 |
| atctcgttgg | agccctggag | gagcagggcc | ttcttgaggt | tgccagtctg | ctggtccatg | 300 |
| taggccacgc | tgtttttgca | gtggtagggt | atgttctggg | agg | | 343 |

<210> 419

<211> 255

<212> DNA

<213> Homo sapien

<400> 419

| | | | | | | |
|-------------|------------|-------------|------------|------------|------------|-----|
| cctagcaaga | gaatcaccaa | atthtatggag | agttaacagg | ggtttaacag | gaaggaagtg | 60 |
| ccttttagtaa | gttctcaagc | cagaggctgg | aggcagcagc | taaatcagag | gacagcatcc | 120 |
| tcagtgaag | tgagccattc | ggggtggcat | gtcactccag | gaataaacac | aacttagaaa | 180 |
| caaagtattt | cgtaggatag | cacagtgcac | tggtgcactg | tgaacctgag | gccactgtgt | 240 |
| caaactgtgc | actgg | | | | | 255 |

<210> 420

<211> 261

<212> DNA

<213> Homo sapien

<400> 420

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cttctgatga | taaccaaccc | ctagctacca | ctctgtattc | atcaggggag | gggtataaac | 60 |
| cccacatgca | agaagaaccc | ttgccccag | tgtcaaatgg | gatggggatg | ctagagttat | 120 |
| agtaaagggg | aaaccctatg | taagctgtta | acagagttca | caggggtagg | gataaccctt | 180 |
| gttctccagc | tcccaaatgt | gctcactttc | ccagcttctt | catccgttca | tcaatgctgg | 240 |
| caaagttccc | ctcaactgtg | g | | | | 261 |

<210> 421

<211> 179

<212> DNA

<213> Homo sapien

<400> 421

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccttcctgtt | gttgtttcaa | atgctgcttg | atttctcgta | acagatctgc | atctatgtaa | 60 |
| tacctttctt | cagatctgac | tgctccaaaa | tgattctgca | tcttgatttg | agacatcaat | 120 |

tcatttagtc ggcccttgaa ctgagtaggt gcatttagtt caccctgaat cgtatccag 179

<210> 422
 <211> 424
 <212> DNA
 <213> Homo sapien

<400> 422
 cgaggtccaa atctgatctg cagatgcaga agattcgaca gaagctgcag actaaacagg 60
 ctgccatgga gaggtctgga aaagctaagc aactgcgagc acttaggaaa tacgggaaga 120
 aggtgcaaac ggaggttctt cagaagaggc agcaggagaa agcccatatg atgaatgcta 180
 ttaagaaata tcagaaaggc ttctctgata aactggattt ccttgaggga gatcagaaac 240
 ctctggcaca gcacaagaag gcaggagcca aaggccagca gatgaggaag gggcccagtg 300
 ctaaacgacg gtataaaaac cagaagtttg gttttggtgg aaagaagaaa ggctcaaagt 360
 ggaacactcg ggagagctat gatgatgtat ctagcttccg ggccaagaca gctcatggca 420
 gagg 424

<210> 423
 <211> 256
 <212> DNA
 <213> Homo sapien

<400> 423
 ctgtggccta gggctacctc aagactcacc tcatccttac cgcacattta aggcgccatt 60
 gcttttggga gactggaaaa gggaagggtg ctgaaggctg tcaggattct tcaaggagaa 120
 tgaatactgg gaatcaagac aagactatac cttatccata ggcgaggtg cacaggggga 180
 ggccataaag atcaaacatg catggatggg tcctcacgca gacacacca cagaaggaca 240
 ctagcctgtg cacgcg 256

<210> 424
 <211> 330
 <212> DNA
 <213> Homo sapien

<400> 424
 ccagccgcat gggagtggag gcagtcacg ccttgctaga ggccaccccg gacacccag 60
 cttgcgtcgt gtcactgaac gggaaccacg ccgtgcgcct gccgctgatg gagtgcgtgc 120
 agatgactca ggatgtgcag aaggcgatgg acgagaggag atttcaagat gcggttcgac 180
 tccgagggag gagctttgcg ggcaacctga acacctaaa gcgacttgcc atcaagctgc 240
 cggatgatca gatcccaaag accaattgca acgtagctgt catcaacgtg ggggcacccg 300
 cggctgggat gaacgcggcc gtacgctcag 330

<210> 425
 <211> 333
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(333)
 <223> n = A,T,C or G

<400> 425
 ctgctccatg gnetcaaagt cagcaccacc cacaccaca atgatcactg acatgggcag 60

```

gttcgaggca cgcaccacag cctcacgtgt ggcttcacaca tccgtcacag caccatcagt      120
cagnagaaac agnatgaagt attgngagggc antcccctga tgtgcagcct gggctgcaaa      180
cctggacctg cccgggcggc cgctcgaaaag ggcgaattcc agcacactgg cggccgttac      240
tagnggatnc aganctcggg acnaagcttg gcagtaatca tggtcatagc tgtttcctgt      300
gagcggttgg gatgaacgcg gccgtacgct cat                                     333

```

```

<210> 426
<211> 411
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(411)
<223> n = A,T,C or G

```

```

<400> 426
gggtgttcat catgaggatt gcttctgcca tggagctgat ggacgtgggc aggttgctga      60
gaaggtgggg tggaagtga tgcggggggg ggggtgagtgc cctggtcttg ttcatagggg      120
agcctttccc tagcagtga acgctgtggg cattttctct agcatattcc cttgggaagt      180
ctagatttgc tattaatctg gctgagaatc taagttctgt gccttagaga cagtttgcac      240
tttcccatat tgtgcctggg acagccatat gatttttttt cccaccaaac aagtatgcaa      300
acagaaacca gttcaaaggg ggatgggtga aaagatgagg cagtanaaat gcctttgaat      360
ggttttctgt agctaattct ctttaaattt tgtcctgctt tttttcttta t                                     411

```

```

<210> 427
<211> 450
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G

```

```

<400> 427
acgtgtacaa gtttgaactg gatacctctg aaagaaagat tgaatttgac tctgcctctg      60
gcacctacac tctctactta atcattggag atgccacttt gaagaaccca atcctctgga      120
atgtggctga tgtggnatc aagttccctg aggaagaagc tccctcgact gtcttgctcc      180
agaacctttt cactccaaaa caggaaattc agcacctgtt ccgcgagcct gagaagaggc      240
ccccaccgt ggtgtccaat acattcactg ccctgatcct ctgcgcgttg cttctgctct      300
tcgctctgtg gatccggatt ggtgccaatg tctccaactt cacttttgct cctagcacga      360
ttatatttca cctgggacat gctgctatgc tgggactcat gtatgtctac tggactcagc      420
tcaacatggt ccagaccttg aagtacctgg                                     450

```

```

<210> 428
<211> 377
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(377)
<223> n = A,T,C or G

```

<400> 428

```

cagggctata gtgcgctatg ttgatctggt gttcatgcta agttccgcat caatatggtg      60
acttcttggg agtggggggac caccagggttg cctaaggagg ggtgaacctg cctacgttgg    120
aaatagagct ggncaaaaact cctgtgctca tcagtagtag aattgcacct gtgaatagcc      180
nccgccctcc agcatgggca acataacaag accctgcctc ttaaagataa aaattggaaa      240
acactngtag gaaaaaaagg gtgnttggtc taaataaatn tggattgggn ataaatgacn      300
caaaactatc atgaatttga aagcntttct aatttcttga aagtctgaaa aaagttaaan      360
cncaatttta tctnaaa

```

<210> 429

<211> 206

<212> DNA

<213> Homo sapien

<400> 429

```

gttgctcctc caaagaaggt tggcttcaag gccgtgtcca gggacccacg agcagaggca      60
ctgggggggca agggatctcc aaggggggcaa gggatcccta aagggggtag ctcacagggtg    120
aggggggttta gggccctct agggagcgcc tgaggccata cattcaagag tgtccctggg      180
gaggcccgag gaagagccag gactgg

```

<210> 430

<211> 473

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(473)

<223> n = A,T,C or G

<400> 430

```

ccttatttnt cttgtccttt cgtacaggga ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta      120
atagcggctg caccatcggg atgtcctgat ccaacatcga ggtcgtaaac cctattgttg      180
atatggactc tagaatagga ttgcgctggt atccctaggg taacttggtc cgttgggtcaa      240
gttattggat caattgagta tagtagttcg ctttgactgg tgaagtotta gcatgtactg      300
ctcggagggt gggttctgct ccgaggtcnc cccanccgaa atttttaatg cagggttggg      360
agntnaggac ctgtgggttt gttaggtact ggggtgcatta ataaattaaa gctccatagg      420
gtcttctcgt cttgctgtgt tatgcccnc tcttcacggg caggtcaatt tca

```

<210> 431

<211> 215

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(215)

<223> n = A,T,C or G

<400> 431

```

cctgtatnaa gctanaaaaa gactaccagc ccgggatcac cttcatcgtg gtgcagaaga      60
ggcaccacac ccggctcttc tgcactgaca agaacgagcg ggttgggaaa agtggaaaca      120

```

ttccagcagg cactgactgtg gacacgaaaa tcacccaccc caccgagttc gacttctacc 180
 tgtgtagtca cgctggcatc caggggacaa gcagg 215

<210> 432
 <211> 391
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(391)
 <223> n = A,T,C or G

<400> 432
 ccagcactgc cacaaacttt ttcagggcca ccaggcgtg cccttcagg accgggaacc 60
 tgcccacttc tatccgcagg atgtagtgc gtgcagattc caggtcagcc atgtagatcc 120
 tggagcgatc tgccaatttc caaacagtgg gagctatctt gttagcagt gttgggtgcaa 180
 ctgtggtctg ggcagcctcc ctggtgagcc cagagagtct ctgcaggtaa gcggtataga 240
 aggacctgga ttccatgagc acggggactc gggagacgga gccattccgg aacagcaggt 300
 agcaagaggg gaagtcgggtg acaccaaact ttctcaccac attggcctct gtgttcagca 360
 ccctgcgcac cgccacncct ttgtgctggg a 391

<210> 433
 <211> 420
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(420)
 <223> n = A,T,C or G

<400> 433
 ctgtagcttc tgtgggactt ccactgctca ggcgtcaggc tcagatagct gctggctgcg 60
 tacttgttgt tgctttgttt ggaggggtgt gtggtctcca ctcccgctt gacggggctg 120
 ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
 agtgtggcct tggttgcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg 240
 gcagccttg gctgacgtag gacggttagt ttggnccctc cgccgaatgc cgcanttcta 300
 ctgtcccaca cctgacagta atagtcanc tcattctcgg cttgggctct gctgatggtc 360
 aggggtggccc gtgntccccg agttggagcc agggaatcnc tcagggatcc canagggccn 420

<210> 434
 <211> 239
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(239)
 <223> n = A,T,C or G

<400> 434
 ccaaccanga gagaaggat cgcttgggtg ccagggccca ccaggagctc caggccact 60
 tgggattgct gggatcactg gagcacgggg tcttgtagga ccaccaggca tgccagggtc 120

taggggaagc cctggccctc aggggtgtcaa ggggtgaaagt gggaaaccag gagctaacgg 180
tctcagtgga gaacgtggnc cccctggacc ccagggtctt cctgggtctgg ctggtnacg 239

<210> 435
<211> 415
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(415)
<223> n = A,T,C or G

<400> 435
ctgtccaatg gcaacaggac cctcactcta ttcaatgtca caagaaatga cgcaagagcc 60
tatgtatgtg gaatccanaa ctcagtgagt gcaaaccgca gtgaccagc caccctggat 120
gtcctctatg ggccggacac ccccatcatt tccccccag actcgtctta cctttcggga 180
gcaaacctca acctctctg ccaactcggc tctaaccat cccncanta ttcttggcgt 240
atcaatggga taccgcagca acacacacaa gttctnttta tcgccaaaat cagcccaaat 300
aataacggga cctatgcctg tttagggntn taacttggnt actggccgca anaattccat 360
agtcaagagc atcacagnct ctgcatntgg aacttctcct ggctntcaga cctgn 415

<210> 436
<211> 152
<212> DNA
<213> Homo sapien

<400> 436
ccaggattga caggccatcc attcacagcc aggagatgct gggccagtc ctccaagagg 60
tctccgtcat ggcagtgatg aaaacctaac aggggtggccc cctgtgccag ctcaggtgac 120
tggagcccga gggcctgaca ggttcccagc ag 152

<210> 437
<211> 174
<212> DNA
<213> Homo sapien

<400> 437
ccagggtactg gcacatcatg ctctggatgg ggggtgggtg gtctgtgaag cagagaaaca 60
ggaaattgtc gtagtcagta tcgagcagct gtggcctcgt tcgccaccgt atagttgatc 120
ttgaacttct ttggattctc agtcttctct ccaaggacct tcttctcaac acag 174

<210> 438
<211> 485
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(485)
<223> n = A,T,C or G

<400> 438
ccacggccct ctcggccctc tcgctgggag cggagcagcg aacagaatcc atcattcacc 60

<400> 441

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccacacagan | tcaccaagcc | acagacttgt | cttccacaag | cacgttctta | tcttagccac | 60 |
| gaagtgacca | agccacacgt | actaaagggt | gaactcaaag | atatgtacag | ggtattaaac | 120 |
| aaataccaag | gggaacagtt | aacttcaata | caaggtcgaa | atcagcaaca | agttctacaa | 180 |
| tccagnctg | atatcagata | caagcttcaa | ggacaatttc | ttttcgaagg | cttattccag | 240 |
| tttcgngagg | ctagcatgag | gtgtgtgcat | ttgccagggg | caaatttcta | ttctcaatta | 300 |
| acccatgcag | caaagtctac | ncatggtgcn | gagtcggttt | agaagcattt | gcggtggacg | 360 |
| atggaggggc | ccgactcgtc | ttactcctgc | ttgctaatac | acnngngctg | gaaggnggac | 420 |
| agtgaggcca | cggatggagc | caccnatcca | cacogagtnc | ttgcgctctg | ggggtgcgat | 480 |
| natnttgatc | ttcatggtgc | tgggc | | | | 505 |

<210> 442

<211> 386

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(386)

<223> n = A,T,C or G

<400> 442

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| cgccaggtga | tacctccgcc | ggtgacccag | gggctctgcg | acacaaggag | tctgcatgtc | 60 |
| taagtgctag | acatgctcag | ctttgtggat | acgcggactt | tgttgctgct | tgcagtaacc | 120 |
| ttatgcctag | caacatgccca | atctttacaa | gaggaaaccg | taagaaaggg | cccagccgga | 180 |
| gatagaggac | cacgtggaga | aaggggtcca | ccaggccccc | caggcagaga | tgggtgaagat | 240 |
| ggtcccacag | gccctcctgg | tccacctggt | cctcctggcc | cccctggtct | cgatgggaac | 300 |
| tttgctgctc | agtatgatgg | aaaaggaggg | nggacttggc | cctggaccaa | tgggcttaat | 360 |
| gggacctana | ggcccacctg | gtgcag | | | | 386 |

<210> 443

<211> 404

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(404)

<223> n = A,T,C or G

<400> 443

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| cctccctctc | agagcttgcc | ccagggactc | tctggccctc | agggttcaat | gtattctgac | 60 |
| caaggccaag | ctttcctggg | gctcagggaa | aatcacactt | tgctaccoga | agctgtatcc | 120 |
| cctcagatgc | caggaaggcc | gtgatcatct | gactccaccc | tcttgagaca | cattctctcc | 180 |
| ctgactgtcc | tgttctaagt | cagcggagca | ccttaggatg | gaggggtgga | ggcgaggcca | 240 |
| ngatgcagcc | tctgtgaaca | ggtgcctgga | ggctgggaaa | tgaccctgag | agggcaggac | 300 |
| acagcnaccg | ngggcttaag | gtgagggngg | agagcaagnt | tggcccaactt | tacaattcta | 360 |
| gntcagagcc | anccctaac | atggngggca | tttattcatt | tcgg | | 404 |

<210> 444

<211> 318

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(318)
 <223> n = A,T,C or G

<400> 444
 catgggctat agtgcgctat gttgatctgg tgttcatgct aagttccgca tcaatatngc 60
 gacttcttng gagtggggga ccaccangtt gcctaaggag ggggtgaacct gcctacgttg 120
 gaaatagagc tgggtcaaac tcctgtgctc atcagtagta gaattgcacc tgtgaatagc 180
 caccgccctc cagcntgggc aacatagcaa gacctgcct cttaagataa aaattggaaa 240
 aacttggtan gaaaaaaagg ctgtttgggtc taaanaagtc tggatngggg ataatgaca 300
 cnaanctatc atgactnt 318

<210> 445
 <211> 418
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(418)
 <223> n = A,T,C or G

<400> 445
 ccagtccaac ctgctcctca ttattgtata aatgagcaga atcaatatgg cggaagccag 60
 cttcaattgc caatttggtg gcctctaaag ctttactttt aggaacctct gcaggcgcat 120
 aggtgccaaa tcccaggaca ggcataagt gaccatcatt cagcttcaca cactgatatt 180
 tcgaatccat ttctgtcact agcctggctg gcaaagtgtt ctttcttcct ccctcacagg 240
 ctataagagc aatgagctgg caacgcccct gagcacactg tctgctgntt aaccaatggc 300
 atgtgagagg agggacagag gcagtcttac acaagctgtg ataaaaattg catncagttc 360
 aaccagtttc ttacnttatt ctaatngna ggaagtgtgn gaagagcaca aagtcaga 418

<210> 446
 <211> 361
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(361)
 <223> n = A,T,C or G

<400> 446
 ctgtccaatn acaacaggac cctcactcta ctcagtgtca caaggaatga tgtaggaccc 60
 tatgagtgtg gaatccanaa cgaattaant gttgaccaca gcgacccagt catcctgaat 120
 gtcctctatg gcccagacga cccaccntt tccccctcat acacctatta ccgccagggg 180
 gtgaacctca gcntctcctg ncatgcagcc tctaaccacac ctgcacagta tccttggtg 240
 attgatggga acntccagna acacnacaca agagctcttt atctccancn tnactganaa 300
 gaacagcgcg actctatncc ttccaggggg ggggggtggg gnntgnggac cttncggggc 360
 c 361

<210> 447
 <211> 321

<220>
 <221> misc_feature
 <222> (1)...(328)
 <223> n = A,T,C or G

<400> 450
 ctggcaattt tgagctgccg gttatacacc aaaatgttct gttcagtacc tagctctgct 60
 cttttatatt gctttaaatt tttaaagaaa ttatatgtca tggatgtggt tatttgtgca 120
 tattttttta caatgcccaa tctgtatgaa taatgtaaac ttcgattttt ttttaaaaaa 180
 attagatttt agctggagct tttgactaat gttaaagtaaa tgccaaacta ccgacttgat 240
 ngggatgttt ttgtaangtt aatttttctaa gactttttca catccaaagt gatgctttgc 300
 tttgggtttt aactgtttca acntnggn 328

<210> 451
 <211> 209
 <212> DNA
 <213> Homo sapien

<400> 451
 ctgccttggt tcaacagaca tgcaaagatc ctaggagaca gtcccatag accttcagac 60
 attaaaaagg gagccgtaca gtttgtttga agcacttcgt cttaccatt tatgcagggg 120
 ccccaggaaa cttacacaca gccagaatga gggtcccaaa ggacttacat taattatggc 180
 tcttgcttcc tttcacaaat gagctgagg 209

<210> 452
 <211> 457
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(457)
 <223> n = A,T,C or G

<400> 452
 ctgtctantc ccttcaagag ctgtttatag aagcttgaga atggggtaaa aatttctgct 60
 agcaaatca agttcttttt gaaattttat cagtaatcca gaatttagta gtocatgcct 120
 tctcactcag catttagaaa taaaaatgtg gtttcttaaa cgtatatcct ttcattgata 180
 tttccacatt tttgtgcttg gatataagat gtatttcttg tagtgaagtt gttttgtaat 240
 ctactttgta tacattctaa ttatattatt tttctatgta ttttaaagn atatggctgt 300
 ttaattcttg aagcattttg ggcttaagat tgccagcacc acacatcaga tgcagtcatt 360
 gttgctatca gtgtggaatc tgatagagtc tngactccgg ccacttggag ttgtgnactc 420
 caaagctaag gacagtgatg aggaagatgg catgtgg 457

<210> 453
 <211> 277
 <212> DNA
 <213> Homo sapien

<400> 453
 ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt 60
 agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
 atttcttgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
 gcatacagga ctaggaagca gataaggaaa atgactacga gggcgtgatc atgaaagggtg 240

ataagctctt ctatgatagg ggaagtagcg tcttgta

277

<210> 454
<211> 198
<212> DNA
<213> Homo sapien

<400> 454
gttaaaagat agtaggggga tgatgcta atcaggctg tgggtggtg tgttgattca 60
aattatgtgt tttttggaga gtcattgcag tggtagta ataatgttg ggacgattag 120
ttttagcatt ggagtaggtt taggttatgt acgtagtcta ggccatatgt gttggagatt 180
gagactagta gggctagg 198

<210> 455
<211> 608
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(608)
<223> n = A,T,C or G

<400> 455
ctgagcaagc taaggaccag gggcaactag accctaataa tnggtacttt tgaaaatgat 60
acaaactacc ttggttgtaa gaagtgcagg ttgaacactt taggagaaca gtcttcaaac 120
tggcaattca aaatttccca ttatatgtga ataaaattgg aaggatgtta aatgtccatg 180
gaaagttact cttgttaagt aggatgcctt atactgaggc tttanaatga aagtacactt 240
cacaaatgga atagtgaaca taaattacca gaagtcaaga taatagtcac actagtaagg 300
taagcaaggc aaattccctt atacacaaaa attattttga tgaccttttt caataatgaa 360
tctgaaatga agtggtttta aaagctccct aaacacaaaa cgaacataaa actgcttaat 420
aacttttagag ctcatgta atctcttgctg aaacacagta ctgaaattac cagcgaaatg 480
atggaatatc tttaaagcag gncactcngt ataactctgga ataatttcac ttgctaactt 540
ttaagaagta ttctctggac tataaatcnt gggcaaatag acttccactt tattattacc 600
ccaaatta 608

<210> 456
<211> 467
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(467)
<223> n = A,T,C or G

<400> 456
cctggacctg tgtaaacctt caaacactct tttttacatt aggtcgtgaa gttaaatttt 60
ttactgtttc tgtgctacag actcttcaaa gggaaatagt taagtcaatt tcaaagaaaa 120
tgaccagcac attttttaaaa cattagaaat gatttgactt tgactatcta ctgccaaaaa 180
aagggttaagg aatttgtaat gagaagctaa aaactttaag gaattttaag gaactcaaaa 240
caaaaactca ttaaagttaa ttaaagttaa ttctacaaat aaagcctctt aatacatttc 300
tataatagtc acttaagact taaattcaaa cactagcaaa ccacaaaatc agactgtntg 360
actgacatcc aaaagataaa tataaatcaa aatccgaccc cagcattagc caaggggtag 420

gtgttcctct tgaggaaggc aggaattcct cttctgccac ctgttg

467

<210> 457
<211> 183
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(183)
<223> n = A,T,C or G

<400> 457
ccaaattttt tacttttaaac actgaaaaca gaggaagtta ataaaaattt taacctataa 60
agtccccctgg ttgttagtca ttaacagcag attgtcagat aagactggta aaatgatggc 120
tgctaagcat ttgatgatcc aggcgcagga tgatcaaact gcagcagatc atgcacgtga 180
cag 183

<210> 458
<211> 445
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(445)
<223> n = A,T,C or G

<400> 458
gaaaaatata aagccaaaaa ttggataaaa tagcactgaa aaaatgagga aattattggt 60
aaccaatttta ttttaaaagc ccatcaattt aatttctggt ggtgcagaag ttagaaggta 120
aagcttgaga agatgagggt gtttacgtag accagaacca atttagaaga atacttgaag 180
ctagaagggg aagttgggta aaaatcacat caaaaagcta ctaaaaggac tgggtgaatt 240
taaaaaaaac taaggcagaa ggtttttgga agagttagaa gaatttggaa ggccttaaat 300
atagtagctt agtttgaaaa atngnaagga ctttcgtaac ggaagtaatt caagatcaag 360
agtaattacc ancttaatgt ttttggcntt ggactntgag ttaagattat tttttaaatc 420
ctgaggacta ncattaatgg gacag 445

<210> 459
<211> 426
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(426)
<223> n = A,T,C or G

<400> 459
cctatgatan cttctctagc tatcactc caatcagcaa aaaatgagaa aatgttgaga 60
aatagaagat aattcctcat ttaaggccac cttctagaat ttgtgcttaa gattctgctt 120
tcttctcatg ggccagcact tcggcaactg gcaaaaatta ggtgtacagg gatctaggta 180
atactgttta tttagcaat aatatattgt gctaactgtc aggcaccta ttactgagaa 240
ataagggaaa atgagtgtaa agtacaacta agagtctcgg cgacagggaa aaataccatc 300

agttaaatat ccatagtcct agagcattta tgtaaaactg caatntgaat cctgcaatac 360
 atnttggctt tttccctcag tgataccatg tgaggggaagn ngctctgtca aggcggggccg 420
 gataga 426

<210> 460
 <211> 348
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(348)
 <223> n = A,T,C or G

<400> 460
 ccaaatttta aaatgttatt tttcatatca tttataacct tgtcacaatc cacttaaaga 60
 agtttgggta tatttctactg aaaattttct tccagagtag gttttttttc gtgggttggg 120
 gggtaacttt actacaatta gtaagtntgg tgcagaatth catgcaaag aggagtgag 180
 cagngtgata atttaaacat atntaaacaa aaacaaaaaa aatgaatgca caaacttgct 240
 gctgcttaga tctactgcagc ttctaggacc cggtttcttt tactgatnta aaancaaaac 300
 aaaaaaanta annacnttgt gcttgaaatg aancttggtt tttntna 348

<210> 461
 <211> 378
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(378)
 <223> n = A,T,C or G

<400> 461
 ccactaagac agaacggaat ctagtagaag tgcaccaatg cttcagtccc tctactcag 60
 catgggtgagc agtgggtcaat ctgtgccctg tggagatgatg ggcagataat tctggcatgt 120
 gtaaataata ataaataatt cacttggtgc aggcagtagt tctatgaatt aaaacctagt 180
 gtgtacacag tgcctacatg tgttacagcc ccacagtagg aatctacacc aaaatattta 240
 ttagaaggaa tttgggtccgt actacatcac gctttccgga gggtaaaaaa taaagtccat 300
 ctatagacat ttcaccacag acccagagac tgagtctggc taaaacctgc aaaatgtcta 360
 taacaaaagn ggatggct 378

<210> 462
 <211> 197
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(197)
 <223> n = A,T,C or G

<400> 462
 gcgaggtcca cactattaaa agctgttggg taattgaagg tgatataaaa tgactgtcnt 60
 catttggagt gngcagcaca nttacttcat gttgctcang tttanaacaa tntccctgn 120

aagttctcac acagatnggn agaaatcata cctantntng gtnaatcact atggcagccg 180
tngaagaatn taagaga 197

<210> 463
<211> 279
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(279)
<223> n = A,T,C or G

<400> 463
cataagtgat gangaggnaa aatcantnaa taagcctaca acntagaata cattaaaact 60
tgcacatata catgttcaca gcatgtatac aatgataatc cctacggttt aaccaagtta 120
tggttccctt ctacagcaga cacaaaacca aggtgaacta ggtnggcaga tgtanaggga 180
ataccaaaaa aagggtaatn ngntcactga ttctgaagna tntgactgan catactgagc 240
ttctgnactt tggaatgca tnnaggnaac aatatcttg 279

<210> 464
<211> 552
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(552)
<223> n = A,T,C or G

<400> 464
gatgggttga taggtgcagc aaaccaccct ggcgcatgtt taccaatgta acaaacctgc 60
acatcctgca caggtactcc aaaactaaaa gtaaaaaaat ctaaaagaaa aaagaaaaag 120
aattaaacc aaatcactt ccccatctgg acttgattta gatgaaaagc ttctggactt 180
tgagctgatg ctatagtggg ttgaaaattt tggggtcttc agaaggggat gaggatatat 240
tgcagtggag agcaacatga atcatngaga gccagagtat agagagnggt gggtagactg 300
taggagagcc ctcaatgatc ccggctgtct tgtattcgcg ttgcacttac ttgtataata 360
tggcagatgg gatgtgatgt cactttcaag attangttat aaatagacta tggcttcaat 420
cagaggggtt tcttctctgt ctanctctct tttgggtagn ttcatctctga gagaaagcca 480
nacctcngcc gcnaccacg ctaaggggcg anttcacagn cactggcggc cngttactag 540
tggatccgng ct 552

<210> 465
<211> 444
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(444)
<223> n = A,T,C or G

<400> 465
ccactcttgg tagaaacctt gaaactttca ccttgctggg ctttagcaaa gtttcctttt 60

```

acagttctgt ttatgagctt cagctactga taaagcactt cctgaacttc tctattatca 120
tagngaccct ctgaataacc tgagtgactg gctcggcaat tcgctttata accattctta 180
ttcccaaagt tggagcacat aaacatttag atgtcttttc ctgtaaaata ttctagacat 240
ttaccctaac tctagttcaa catatactca acttgcaactg tatactctccc tgcttttttg 300
agacagagaa gaaattcagg aggtgnoccc tctccagagt ttctctgttg gaaagcagcn 360
atcaagaanc ctttaaaaaa ttggtgtnaa gctntgcnc ctgcagaaat gcntngcccc 420
acattattct tctggggnaa agna 444

```

```

<210> 466
<211> 381
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(381)
<223> n = A,T,C or G

```

```

<400> 466
cctactatgg gtgttaattt tttactctct ctacaagggt ttttcctagt gtccaaagag 60
ctgttcctct ttggactaac agttaaat tacaaggggat ttagagggtt ctgtgggcaa 120
atttaaagtt gaactaagat tctatcttgg acaaccagct atcaccaggt tcggtagggt 180
tgctgcctct acctataaat cttcccacta ttttgctaca tagacgggtg tgctctttta 240
gctgttctta ggtagctcgt ctggnttcgg gggctcttagc tttggctctc cttgcaaagt 300
tatttctagt taattcatta tgcannaggt ataggggnta gtccttgcta tattatgctt 360
ggttataatt tttcatcttt c 381

```

```

<210> 467
<211> 95
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(95)
<223> n = A,T,C or G

```

```

<400> 467
cctatanatt ntggnttgta tactgggtcc tgaaaaccct cttggngctc tgtttttaag 60
gagctgaanc caanganccg caataataat acttt 95

```

```

<210> 468
<211> 224
<212> DNA
<213> Homo sapien

```

```

<400> 468
cagtgggtct ctgatgcctt gcctgcagca gaaggaggga gcagagatca agaggaagga 60
aaaaatcata tgtacttatt tgaaggtaaa gattattcta aagagcccag taaggaagac 120
agaaaatcat ttgaacaact ggtaaacctt cagaaaaccc ttttggagaa agctagtcaa 180
gagggccgat cactccgaaa taaaggcagt gttctcatcc cagg 224

```

```

<210> 469
<211> 416

```

<213> Homo sapien

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ctgagttcta | gttcaaaagc | tttatcctta | acttcgtcat | gtactatgta | aattctagaa | 60 |
| tagaaaaggg | aaaggtgaaga | ttttggtaac | ctccaaacat | tgaagtagtt | cacagaccca | 120 |
| aagtcagtac | aaattagaat | gtccatccat | aataaaagta | tctataaaat | tacacagaca | 180 |
| cattctacat | agtatttaac | attagagaag | acaaattaca | cagggactga | aataaaatga | 240 |
| aacatctact | ctcccgacaa | atgttgaata | tacctaatca | acccaagttc | agtttatttt | 300 |
| tgcacattgc | tttagagata | taacttggct | gggcacagtg | gctcacacct | gtaatcccaa | 360 |
| cactttggga | gaccaaggcg | qatqqatcac | ttgaggtcag | ttcgagacta | gcctgg | 416 |

<213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| caccttttaaa | ctgtatcaca | aagtctgttg | ctgtggttac | agcctttgtt | tccagtgatg | 60 |
| ttttgtccat | gctttccccc | aacccttaac | aatggttact | caaaagaatg | aaataatgag | 120 |
| tcattcattc | gggaatatgt | taaaatatcc | ctctttatca | ttacatttca | ctgcttagaa | 180 |
| actaggctgt | aattcaaggc | aacagttaag | tctgagaact | gttaaaaaaa | tctttgattt | 240 |
| tttttcattt | ttaagaaaaa | cctgcctatt | taattgttca | gacttgtaag | aggttcttca | 300 |
| attacatcct | ttttggttaa | tgtattattt | ctggaacaag | tagataaaat | tctacgcagt | 360 |
| aaacataata | aaaatc | | | | | 376 |

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ggcttcgtat | aatggttctt | ttgtcacccc | tgatcgacga | tttcgctacc | cgtacaactc | 60 |
| tgacaaggga | acgaaatgct | tctgtgtatt | cacctagtgg | tctgtgaac | agaagaacaa | 120 |
| caactccacc | ggatagtggg | gtactgtttg | aagggttagg | catttcaaca | agacctagag | 180 |
| atgttgaaat | tcctcagttt | atgagacaga | ttgcagtaag | gaggccaact | acggcagatg | 240 |
| aaagatcttt | gcggaaaatt | caagaacaag | atattattaa | ttttagacga | actctttacc | 300 |
| qtgctggtgc | tcgagttaga | aatattgaag | atggtggccg | ctacagggat | atttcag | 357 |

<213> Homo sapien

<223> n = A, T, C or G

| | | | | | | |
|------------|-------------|------------|------------|-------------|------------|-----|
| cngagatgac | atttacaatc | tcttgaaang | cagcagatgg | cactctgggtg | cttcttatga | 60 |
| agcaacatgc | ttgaaatcaa | gggccaaaca | ttgttgtagg | aaagcaaaat | atacctctaa | 120 |
| cacctacgtt | taccaaiaaaa | gctgacatct | caaactctga | gttggttgaga | ctcaaatttc | 180 |
| tcatcccca | aqaaqcctat | tacggtagtg | tgntggatgc | tttttgtatc | tctgataggc | 240 |

```
<210> 473
<211> 264
<212> DNA
<213> Homo sapien
```

```
<210> 474
<211> 165
<212> DNA
<213> Homo sapien
```

```
<210> 475
<211> 417
<212> DNA
<213> Homo sapien
```

```
<220>  
<221> misc_feature  
<222> (1)...(417)  
<223> n = A,T,C or G
```

```
<210> 476
<211> 321
<212> DNA
<213> Homo sapien
```

<220>


```

gnnttccaaa ttcttctaac tcttccaaaa gccttctgcc ttagtttttt tttaaattaca      60
ccagtccttt tagtagcttt ttgatgtgat ttttaaccaa cttcccttc tagcttcaag      120
tattcttcta aattggtoct ggtctacgta aacacctca tcttctcaag ctttaccttc      180
taacttctgc accaccagaa attaaattga tgggctttta aaataaattg gttaccaata      240
atttcctcat tttttcagtg ctattttatc caatttttgg ctttatattt ttctatcttc      300
tatacttctc caatacttgt cttagcttgt ttttcatttt ctatctgaaa ctcttgacaa      360
tatcttctaa tttccctatc ttctctatct ttttcttcgc cttcccgta tctgcttcc      420
agntttccac ttcaaacttc tatcttctcc aaattgttca tctaccact cccaataatc      480
tttccatttt cgtgtagcac ctggncag      508

```

```

<210> 480
<211> 81
<212> DNA
<213> Homo sapien

```

```

<400> 480
ggtgcccttt tcctaacact cacaacaaaa ctaactaata ctaacatctc agacgctcag      60
gaaatagata aggaaaatga c      81

```

```

<210> 481
<211> 306
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (306)
<223> n = A,T,C or G

```

```

<400> 481
tcgccttcgg ccgccgggca ggtaggggn acaagacgct acttccccta tcatagaaga      60
gcttatcacc tttcatgac acgccctcat agtcattttc cttatctgct tcttagtcct      120
gtatgccctt ttctaacac tcacaacaaa actaactaat actaacatct cagacgctca      180
gggaatagaa accgtctgaa ctatcctgcc cgccatcctc ctagtcctca tcgcctccc      240
atccctacgc atcctttaca taacagacga ggtcaacgat cctccctta ccatcaaatc      300
aattgg      306

```

```

<210> 482
<211> 582
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (582)
<223> n = A,T,C or G

```

```

<400> 482
ggggggaaca gtcattatac attatttaga ctcatctcct cttccagtgc ccttatgatt      60
atttcttacc tttaccattg atcttaaaact gngcaggcta aaaagaggaa ccagaactcc      120
cttaagcact tttaagacta tttaaaaaat aaagntttgt tggcattgaa gagtaagctg      180
cttaagggac tgaatgaaaa gatagtaccc tttgtggctg tatgaagaga gaaactgaat      240
ttctatccaa gagaccttaa tntagcctat tagggaatta tcttcccaa aagtacaagt      300
aattttgcac tgcaggagaa ggataagtag atttgattta catcacattt tatacacacc      360

```

```
<210> 483
<211> 275
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(275)
<223> n = A,T,C or G
```

```
<210> 484
<211> 434
<212> DNA
<213> Homo sapien
```

```
<210> 485
<211> 291
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(291)
<223> n = A,T,C or G
```

| | | | | | | | |
|------------|------------|------------|------------|------------|------------|--|-----|
| <400> | 485 | | | | | | |
| ncaccactgc | agccctacat | acagttgaaa | aaaaattcca | ttctgttaac | atttgtttta | | 60 |
| taagttttca | cgcaatacac | aaaaaacccc | tctgcacttc | ttgtaaagaa | caaaaaagat | | 120 |
| acacaacagt | taaggcgtaa | gatacacagg | aatagcattc | aaacatggat | gtgggtagag | | 180 |
| aaaggagtag | ctggcgtagt | tacctgcctt | gtttgactga | atccttgatt | tttaatttgg | | 240 |
| cttttcatgg | qccgctcaca | acaccaacgc | tgtgtgaggt | atggtagtca | g | | 291 |

<210> 486
 <211> 274
 <212> DNA
 <213> Homo sapien

<400> 486
 ctgtaatat gtagttgctc cagaatgtca agggcagctt acggagatgt cactggagca 60
 gcacgctcag agacagtga ctagcatttg aatacacaag tccaagtcta ctgtgttgct 120
 aggggtgcag aaccgcgttc tttgtatgag agagggtcaaa gggttgggtt cctgggagaa 180
 attagttttg cattaaagta ggagtagtgc atgttttctt ctgttatccc cctgattgtt 240
 ctgtaactag ttgctctcat ttttaatttca ctgg 274

<210> 487
 <211> 184
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(184)
 <223> n = A,T,C or G

<400> 487
 tggcaccaag attctcagct cacggtacca gcatctgatt gtcggactac ctgctgcttt 60
 ccctgatatt tatacatgat attcgnaaaa tgtaaagaag ctattattca tacagacatc 120
 tagagaagga gngaagnttt taaaaaaata aaaaaatact tatttcaagc tttagctgtg 180
 ttct 184

<210> 488
 <211> 393
 <212> DNA
 <213> Homo sapien

<400> 488
 ctgcattttt attgcgatct gcagatgaac tggaaaatct cattttacaa cagaactggg 60
 acagacgacc accatattca ctgaggtcta aatttgagct ttccactaat gacattttga 120
 tttcccaaca gagatacttc tggctcttact gcacagtctt ttaagagaaa tacttccatt 180
 atgccacatt gtccttgatc cgtaagtgat gtgttaaggt gcttcaaagg aactctgacc 240
 tctgaagtac ttgagctact ttagtatgtc cagcctattg ctttttggtt tagtgtgtca 300
 ccataaatat caggggcata aaaggctatc tattcttaat tcaaggataa aacagaagaa 360
 gcttggtgga taaaacaata gttcaagatc cag 393

<210> 489
 <211> 607
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(607)
 <223> n = A,T,C or G

<400> 489
 gtgcttatgt acttaagggg aactactcta actgggtgaa gagtangatg aagcatccat 60


```
<210> 490
<211> 179
<212> DNA
<213> Homo sapien
```

```

<400> 490
cttctaggaa tactagtata tcgctcacac ctcatatcct cctactatg cctagaagga      60
ataatactat cactgntcat tatagctact ccataaccc tnaacaccca ctccctetta    120
gccaatattg ngcctattgc catactagtc tttgccgcct gcgaagcanc ggtaggacc     179

```

```
<220>  
<221> misc_feature  
<222> (1)...(399)  
<223> n = A,T,C or G
```

```
<210> 492
<211> 482
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(482)
<223> n = A,T,C or G
```

<400> 492
ctccacctta ctaccagaca gccttagcca aaccatttnc ccaaataaag tataggcgat 60
agaaattgaa acctggcgca atagatatag taccgcaagg gaaagatgaa aaattataac 120
caagcataat atagcaagga ctaaccccta taccttctgc ataatgaatt aactagaaat 180
aactttgcaa ggggagccaa agctaagacc cccgaaacca gacgagctac ctaagaacag 240
ctaaaagagc acaccgtct atgtagcaaa atagtgggaa gatttatagg tagaggcgac 300
aaacctaccg agcctggtga tagctggttg tccaagatag aatcttagtt caactttaaa 360
tttgcacaca gaacctcta aatccccttg taaatttaac tgttagtcca aagaggaaca 420
gctctttgga cactaggaaa aaaccttgta gagagagtaa aaaatttaac acccatagta 480
gg 482

<210> 493
<211> 207
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(207)
<223> n = A,T,C or G

<400> 493
cataaatatt atactagcat ttaccatctc acttngngga atgctagtat atcgctcaca 60
cctcatatcc tccctactat gcctagaagg aataatacta tcaactgttca ttatagctac 120
tctcataacc ctcaacaccc actccctctt agccaatatt gtgcctattg ccataactagt 180
ctttgccgcc tgccaagcag cggtagg 207

<210> 494
<211> 283
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(283)
<223> n = A,T,C or G

<400> 494
ccaattgatt tgatggtaag ggagggatcg ttgacctngt ctgttatgta aaggatgcgt 60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaagggtg 240
ataagctctt ctatgatagg ggaagtagcg tctttagtagac cta 283

<210> 495
<211> 590
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(590)
<223> n = A,T,C or G

<400> 495
 tatgtatata attttcttag ttactagcat agagaaatta ctgatttaaa aaaacatttc 60
 aaattctagc atgttgtagg attctattgc cctttctaaa aagtacatct tgcttatccg 120
 atttctaaca aaactattta atttgaagaa gggagaatga atttggataa aaagcaaaaa 180
 tttaaaggta ctcaaattta ggcaaaccat taaagcaatc ttagtttaca gttaattggg 240
 tagaatggtc aacactttct tcagggttagt tcatggagtg gatatgcatt gatagaacaa 300
 cttagagatg cttttacagt tgagaaagct cattatatct gttatcttta agaatcagct 360
 tatttatttc atatgtttgt tctttaagaa gaccaaagag ccctgcaaata gaatgttgat 420
 ttgttttttt gtttgtttaa ttttttga gagataagat ctcactttgt tatgttgccc 480
 aggctggtct caaactctca acttgaagtg atctgccac ctcagcctcc caaagtgggtg 540
 ggattacagg catgagccac cgcacctgga cctgcccggg cggncgctcg 590

<210> 496
 <211> 307
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(307)
 <223> n = A,T,C or G

<400> 496
 ggagattagt atagagaggn anacnttttt tcngnatatt tggtcacatg gataagtggc 60
 gctggcttgc catgattgtg aggggtagga gccaggtagt tagtattagg aggggggnng 120
 ttagggggtc tgaggagaag gttggggaac agctnaatag gttgttngnt gatttggnnta 180
 aaaaacanta gggggatgat nctaataatt antgctgtgg gtggttgn tgattcaaata 240
 tatngctttt ttcggagann catgtcangt ggtagtaaata ataattgttg ggaccattan 300
 ttcttan 307

<210> 497
 <211> 216
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(216)
 <223> n = A,T,C or G

<400> 497
 cattttcttc ttggtttctt cagttaagtc aaannngnac gttcctcttt ccccatatat 60
 tcatatattt ttgctcgtaa gtgtatttct tgagctgttt tcatgttggt tatttctctgt 120
 ctngaaatg gtgttttttt ttgttgtgn tggttttttt tttttttttt aaactnngna 180
 ccncaantt gaaaaaatgn ttntttttcc ctnaca 216

<210> 498
 <211> 375
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature

<222> (1)...(375)

<223> n = A,T,C or G

<400> 498

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| gaatttcctg | gcaccttttc | tcgctagaga | agattnngtg | tgactgggtt | gcctataagc | 60 |
| catatagata | caaactttta | tctctaatac | caagtcttag | agggatatat | taatagatct | 120 |
| aataaattta | ttcttagact | tattgtttca | tgggntagtg | agtctttgct | actggagaca | 180 |
| atacagactt | gtcagttttt | ttaaaaaaaa | aaaatttgcc | aagctancac | attaaaaana | 240 |
| tntcctaagg | ctntcatttt | atgaggatga | ttataaacnt | ttntgngata | aatatcacca | 300 |
| taataaactg | ttaagtacaa | ctgcnggccn | cccttanagn | gaattcctnc | agttanaaat | 360 |
| ttattttttt | gccaa | | | | | 375 |

<210> 499

<211> 215

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(215)

<223> n = A,T,C or G

<400> 499

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccacnaaagc | agaagcttaa | agcatagtag | taaagagggn | aaaaagaagg | acgaaaataa | 60 |
| atcagatgac | aaggatggta | aagaagttga | cagtagtcat | gaaaaggcca | gaggtaatag | 120 |
| ttcactcatg | gaaaagaaat | taagtagaag | gttggtcgaa | aatcggagag | gaagcttgtc | 180 |
| acaaaaaaaa | aaaaaaaaaa | aaaaaaaaat | gtttt | | | 215 |

<210> 500

<211> 489

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(489)

<223> n = A,T,C or G

<400> 500

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccactacgat | aagcaggtag | ctgggttttg | tagtgagntt | gctccttaag | ttacaggaac | 60 |
| tctccttata | atagacactt | catttttcta | gtccatccct | catgaaaaat | gactgaccac | 120 |
| tgctgggcag | caggagggat | gatgaccaac | taattcccaa | acccaggtct | cattggtacc | 180 |
| agccttgggg | aaccacctac | acttgagcca | caattggttt | tgaagtgcac | ttacaaggnt | 240 |
| tgtctacttt | cagttcttta | ctttttacat | gctgacacat | acatacactg | cctaaataga | 300 |
| tctctttcag | aaacaatcct | cagataacgc | atagcaaaat | ggagatggag | acatgatttc | 360 |
| tcctgcaaca | gcttctctaa | ttatacctta | gaaatgttct | cctttttatc | atcaaatctg | 420 |
| ctcaagaagg | gctttttata | gtagaataat | atcagtggat | gaaaacagct | taacatttta | 480 |
| ccatgctta | | | | | | 489 |

<210> 501

<211> 286

<212> DNA

<213> Homo sapien

<400> 501
 aaaaacactc aaacacagcc ttggagggag gagtcagttt taaaagactc ttataaaaagt 60
 aatatactgc tagctctgaa gaatcggagg ctaaaatcat ctcttcaagt cccagggaa 120
 tcccaaagaa ctccagggga aggtgggatg ggccagagag ctctggaagc ttccaggtct 180
 gttgcaagcc tcacctggta cacagtaggc tcttccaggt ctgtcaggaa cccaggagcc 240
 tcccctagca cacagtaggc tcacaaaaag ggagcactgc tgctgg 286

<210> 502
 <211> 168
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(168)
 <223> n = A,T,C or G

<400> 502
 cctatgattg tgggggcaat gaatgaagcg aacagagntt cgttcatttt gggtctcaga 60
 gtttggtata attttttatt tttatgggct ttggtgaggg aggtaagtgg tagtttgtgt 120
 ttaatatatt tagttgggtg atgaggaata gtgtaaggag tatggggg 168

<210> 503
 <211> 173
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(173)
 <223> n = A,T,C or G

<400> 503
 cctttataat aaattaggca aaaggttcag tgcnnngcta tantggacaa catgaaactc 60
 cataaaaatg actggatagg gggactgctt gagacttttc ttttgggcat tactaacaga 120
 attcaaagaa attccaacca cgcttatttt tccaaattct actgaaatga gag 173

<210> 504
 <211> 310
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(310)
 <223> n = A,T,C or G

<400> 504
 tagtattcta tttaaaaatt aagttttggg gtctgtaaaa tatacaggac aatgactttt 60
 ttaaaatgta agttaatacc tcctcctcac ttgtcttaat tgaacttagg tgtttattct 120
 taaagngnga ccttgatgaa aatgttgaga tgggaagtgt tattaggcaa aacttggtat 180
 agattttctca tataactctt aattgaccct tagaatttta acaaccgcgc ctggcccaat 240
 agactgtttt ttagagtant tttaggctct cancaaaatt gaggggaaaa tacagggtgt 300
 tcccattaaa 310

<210> 505
 <211> 530
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(530)
 <223> n = A,T,C or G

<400> 505
 cctcagggaa cttacaatta tggcaaaagg ggaaggggaa gcaagcacct tcttcacaag 60
 gcatcaggag agagagagaa agagagtagg ggaaactacc ctttttaaac catcatatcc 120
 tgtgagaact ccctcagtat tagaagagca tgagggaaac cgctccata atccaatcac 180
 ctcccaccag gaccatccct caatacatgg gggttacaat tcaagatgag gttcgggtgg 240
 ggatacagat ttaaaccata tcagaatggg taatgatatt gttgtatatt accaactata 300
 atcttcttag tggtatagta caataatgta aaaaattgag taaatttggt ttctatatta 360
 ttctgttttt ggaaaacatg tatatagtca gggctgtttg tctcaagaaa atatggtaaa 420
 ctctgctgtt ttggtcactg gtgcctagaa tttggggatg tacattgggt ttgattcaca 480
 tgcacatttc cttctagtgc acagtaacta tttctaacta tttcccnata 530

<210> 506
 <211> 352
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(352)
 <223> n = A,T,C or G

<400> 506
 cttgaacgct ttcttaattg gtggctgctt ttaggcggtg ctatgggtgn taaatTTTTT 60
 actctctcta caaggttttt tcctagtgtc caaagagctg ttctcttttg gactaacagt 120
 taaatttaca aggggattta gagggttctg tgggcaaatt taaagttgaa ctaanattct 180
 atcttgga accagctatc accaggctcg gtaggtttgt cgctctacc tataaatctt 240
 cccactattt tgctacatag acgggtgtgc tcttttagct gttcttaggt agctcgtctg 300
 gtttcggggg tcttagcttt ggctctcctt gcaaanntat ttctagttaa tt 352

<210> 507
 <211> 370
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(370)
 <223> n = A,T,C or G

<400> 507
 cctaactaga tcttatcaga atagggggga agggngtcgg ttcattcetta ttgagtgtta 60
 atgaccctgt aagatgtaat ttcttttatt tcattctgtt acctagaaaa tctatcacag 120
 ccttgtagta ttgattgctc aatctataaa gagctcagtt tacagcatga ctgttagtaa 180

cagggntatt ttaatgagtg actcttcaac acctcagagt ttcactaaat tccaacccat 240
 cagcccagta gtctaacatt aaggggtctta ggaaatgaga acttatcacc ttctcttatac 300
 atgaaaaggt aacctccagg taacccaaaa tagaacttcc tctgtgttcg ttttttatag 360
 aaattactgg 370

<210> 508
 <211> 129
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(129)
 <223> n = A,T,C or G

<400> 508
 ctgttataaag aacaaactta gcaatatata acagttnggt aacaggattt ttgactattc 60
 actttgggag ttatttttaa aaatccactt ttttactgag tcttactaca taccaggcac 120
 tgtacttg 129

<210> 509
 <211> 422
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(422)
 <223> n = A,T,C or G

<400> 509
 ntgggaagtc gtgacatcca tgggaaccca gcgctgtgat gctgggtgttt gngttctccg 60
 cgagaagtga ccattgttgg agcaccatcc agagctagtg accantncag tggacagtta 120
 gtgggagaat caaaaatcct ttccagaatg tctgtttctc actacntgca ccgggngatt 180
 acaggcacca gtgcagngat gattgtactt atttgacaca tactccccgt cntcctggnt 240
 nttgttcctg anaanggtgg gtaaattatc caggaaaaan aatgcacatt gaatggatgt 300
 gagagaccac attgcctctc ccactgcttt ggggagcact ttctgtcat ttctaactta 360
 ccacntgctt ggtgtactat atgtatgttg tgctcatat gttgcaaaga actaangtga 420
 gt 422

<210> 510
 <211> 238
 <212> DNA
 <213> Homo sapien

<400> 510
 ccacctatga attggtgggt tacctactca atggatagca gcacgaggac tgctgtactg 60
 cacaaaaaga agacccaaaag attacagtgg accatgggat acagaagcca gcatggcaga 120
 cagaagaaaa atagtttggg aacatgtaac tctcctaagt ggaagttttg ttgttaggaat 180
 tatagtaatc acaccacatt acttggcctt tcggtaatgt gaaaaaaaaa aaaaatcc 238

<210> 511
 <211> 254
 <212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(254)

<223> n = A,T,C or G

<400> 511

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| ccnattgatt | tgatggtaag | ggaggggatcg | ttgnggctcg | tctgttatgt | aaaggatgcg | 60 |
| tacggatggg | agggcgatga | ggactaggat | gatggcgggc | aggatagttc | agacggtttc | 120 |
| tatttcctga | gcgtctgaga | tgtagtatt | agttagtttt | gttgtaagng | ttaggaaaag | 180 |
| ggcatacagg | actaggaagc | acgataagga | aaatgactat | gagggcgnga | tcatgaaagg | 240 |
| tgataagctc | ttct | | | | | 254 |

<210> 512

<211> 269

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(269)

<223> n = A,T,C or G

<400> 512

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| cctacctgta | aactacagta | ctttatatat | ctatgggntt | aataaaaana | aaatccacaa | 60 |
| atcttaaaaa | ggaactttaa | atgcagggct | atattgaatt | ggnaaaactgc | aacacaaact | 120 |
| ggcgcaacat | aggtaaatga | ataccaatct | cactctatgt | gatgcaagca | tgctaactttc | 180 |
| ccactaattt | aaattacttt | caaccactat | gagccagaat | gcatgctga | accttaaaact | 240 |
| gcactttaaa | aagtaacatc | ttggcctaa | | | | 269 |

<210> 513

<211> 266

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(266)

<223> n = A,T,C or G

<400> 513

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ggaggggggt | tgttaggggg | tcggaggaga | aggntgggga | acagctaaat | aggttggtgt | 60 |
| tgatttggtt | aaaaaatant | agggggatga | tgctaataat | taggctgtgg | gtggttggtg | 120 |
| tgattcaaat | tatgtgnttt | ttggagagnc | atgncantgg | tagtaatata | attggttgaga | 180 |
| cgattagttt | tagcattgga | gtaggtttag | gttatgnacc | gtactctagg | ccatatgtgt | 240 |
| tgganattga | nactagtagg | gctagg | | | | 266 |

<210> 514

<211> 271

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature
 <222> (1)...(271)
 <223> n = A,T,C or G

<400> 514
 acatgcaana aatcgagaat cttaaaaaac annacgaanc tgccttgga nncttactgg 60
 nntangatat ttatnttgcg gctgagatac ttgaacaact tcggatcnga antagacaan 120
 aangggnant tntatactgc nncagagggt acacagntca ttgtattaga gangaacana 180
 tgggtctggt gttcacacat tggggggaan atgggcgttn acangagagg nnganaaacn 240
 anganagcct nctggttng cataanaaaa a 271

<210> 515
 <211> 328
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(328)
 <223> n = A,T,C or G

<400> 515
 ccaatgaggg gcaaagtgag cgncnagaag angttttgac tgaaataaat caaacacaaa 60
 aatntaagtt cacagtgaca gtttaacaa aatccaaaca aactaacaac anaaacaccc 120
 cttgntttgc ctctagtgga aggtgggana acacaanctc gtctataaaa ttgactagta 180
 aaggggaaaa cccggtcatt tncctactct ttccangaaa tatctaatagc aagaaagaac 240
 ttctnctcat tatacngaag gaatttngaa aaatgatgta tttttggaac acctaantga 300
 aatactggaa cctgggcaag ttcaccac 328

<210> 516
 <211> 220
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(220)
 <223> n = A,T,C or G

<400> 516
 nctnagttg aaggacccca tgtacatata gccagggga gcagtactag gntaactaga 60
 aggatctcat ccccatatgt gggctcattt caagtctatg gatgactacc ttcattgntg 120
 tgtgcgagat ggtttcaccc cttgaaaata tgggcacttc ancataanat agcnaaatct 180
 ttataatgat caatncatcc tacctccttt tacatgcatg 220

<210> 517
 <211> 296
 <212> DNA
 <213> Homo sapien

<400> 517
 tgcgatttct tccttggtgt ttgctttggt ctgtgttcaa tccagagagc ttaaattgtc 60
 attatttttg gaagaaaacc tgtatttttg ttagtttaca atattatgaa atttcacttc 120
 aggagaaact gctgggcttc ctgtggcttt gttttcttag tttctttttc cgtgccgtgt 180

```
<210> 518
<211> 299
<212> DNA
<213> Homo sapien
```

<400> 518

```
<210> 519
<211> 464
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(464)
<223> n = A,T,C or G
```

<400> 519

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| gctgcacatc | ggaggaaaaac | tcggtaaagc | agaatgaggt | tgatatgttg | aatgtatttg | 60 |
| attttgaaaa | ggctgggaat | tcagaaccaa | atgaattaaa | aaatgaaagt | gaagtaacaa | 120 |
| ttcagcagga | acgtcaacaa | taccaaagg | ctttggatat | gttattgtcg | gcaccaaagg | 180 |
| atgagaacga | gatattccct | tcaccaactg | aatttttcat | gcctatttat | aaatcaaagc | 240 |
| attcagaagg | ggttataatt | caacaggtga | atgatgaaac | aaatcttgaa | acttcaactt | 300 |
| tggatgaaaa | tcatccaggt | atttcataca | gtttaacaga | tcgggaaact | tctgtgaatg | 360 |
| tcattgaag | tgatagtgc | cctgaaaagg | ttgagatttc | aatggatta | tgtggtctta | 420 |
| acacatcacc | ctcccaatct | gttcagttct | ccagngtcaa | aggc | | 464 |

```
<210> 520
<211> 221
<212> DNA
<213> Homo sapien
```

<400> 520

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| ctgatatctca | cttattttaac | acaagtctct | aatacaatac | aattttatta | attttattcc | 60 |
| acatgcccga | cattagatct | ctagactcat | tcatcctaca | tacctacttt | gtatcctttg | 120 |
| acctacatct | ccctacttcc | tcctccagtc | cccaccccc | accactgggt | gctaaccact | 180 |
| gtttcattcc | ctttttcatt | ctacatatgt | gagatcatgc | t | | 221 |

| | |
|-------|-----|
| <210> | 521 |
| <211> | 312 |
| <212> | DNA |

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| ccagacctgc | agaaaaaactt | agcacagctc | aatctgctgt | tttgatggct | acaggggttta | 60 |
| tttggccaag | atactactt | gtaactattc | caaaaaattg | gagtctgttt | gctgttaatt | 120 |
| tctttgtggg | ggcagcagga | gcctctcagc | tttttcgtat | ttggagatat | aaccaagaac | 180 |
| taaaagctaa | agcacacaaa | taaaagagtt | cctgatcacc | tgaacaatct | agatgtggac | 240 |
| aaaaccattg | ggacctagtt | tattatttgg | ttattgataa | agcaaagcta | actgtgtgtt | 300 |

```

tagaaggcac tgtaactggt agctagttct tgattcaata agaaaaatgc agcaaacttt 360
taataacagt ctctctacat gacttaagga acttatctat ggatattagt aacatttttc 420
taccatttgt ccgtaataaa ccatacttgc tcaaaaaaaaa aaaaaacctt c 471

```

```

<210> 525
<211> 332
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(332)
<223> n = A,T,C or G

```

```

<400> 525
cccnctgta ttccagcctg ggtgacccca tctcanggae gaaaagttac cagatgtcgn 60
gggtaaaggt tggctctcaa gtggcctcat aagttgtctt gcattttaat tcaggggaatt 120
cattggacca ataggttaca ttttcgttcc tttttgtttt tggttcatct gtttaagcagt 180
gggggcctaa ttactgtctc tttgtaaaaa cacattttcc caaagaacac tgaattaccg 240
ttcaaaactgg ttgttgatgg gtaataaggg ctgtttttgc tgccccaaaaa gggcttaaca 300
atttaggcgg atagtttact taaaaaaaaa aa 332

```

```

<210> 526
<211> 440
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(440)
<223> n = A,T,C or G

```

```

<400> 526
ccaggttacc tcccctaaca gatgtggtgt tctgangggg tggttaagtg cccgaggaaa 60
ataggcctta actgttaaca tctacagaga agaaagcatg gtcacactgg caaggagtaa 120
gaagggattg ggtaaaagaa aatgggagag aaaagggaaa aaagtttttg caagacaatt 180
gttcctctgt aagaagctgc agggtgaaaag ctttcctttc ttctattttt gtttttaaatg 240
nctgtctctc tgatcagngg aaaagtgaag atttctagta tctagcacta acgtatgacc 300
caactttgag ggatcacaag ctagaacaag ttgaggattt aaaatcctgg ataattatat 360
acttaaagtt catgagcata aagctcactt gaccatgcag aaatgctggg aagcaggggtg 420
catggcatgg gaatacatct 440

```

```

<210> 527
<211> 124
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(124)
<223> n = A,T,C or G

```

```

<400> 527
tttccatag tctgttgggt gcataaatgn cttcttctga gaagtgtctg ttectatcct 60

```

ttgccccctt tttgaggact taaatgttag acctaagacc ataaaaaccc tagaagaaaa 120
ccta 124

<210> 528
<211> 162
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(162)
<223> n = A,T,C or G

<400> 528
ctgcgggaga aatatgggga caagatgttg cgcangcaga aaggtgaccc acaagtctat 60
gaagaacttt tcagttactc ctgccccaaag ttctgtgcgc ctgtagtgcc caactatgat 120
aatgtgcacc ccaactacca caaagagccc ttctgcagc ag 162

<210> 529
<211> 409
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(409)
<223> n = A,T,C or G

<400> 529
cctttaaaat atagcttata aaatgtatac tatnngccag gagagctcac atttttctgc 60
agttttccag tggacctgcc tatggaatac tgtaaagaaa aatctgcaaa aatattccta 120
gcaattgaat cagtgccttt aaataaaaga agtggagagg ggcttggtta aattattctg 180
acaagttttc ttgctagtgg ttgccaaaat taaggatatt tgaagtgtcc tatcacccaa 240
atttggtttt aagaaaaagc tatattctgn gtctataggg tgaagccac actatctgtg 300
ctgcattctc aatgatacaa tacctatctg gaaactttcc tgttttgcc atgggtgcac 360
aaatctaaaa cattttatca caaagggtac ttgaatttaa atttctttt 409

<210> 530
<211> 325
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(325)
<223> n = A,T,C or G

<400> 530
ccgccagtgt gatggatata tgcagaattc gccctttcna gatttgngcc cgggcaggtc 60
catggctagg attatagata gttgggtggg tggggnaaat gagtgaggca ggagtccgag 120
gagggttagt gtggcaataa aaatgattaa ggatactagt ataagagatc aggttcgtcc 180
tttagtggtg tgtatggcta tcatttggtt tgagggttagt ttgattagtc attgttgggt 240
ggtaattagt cggntgttga tganatattt ggagggtggg atcaatagag ggggaaatag 300
aatgatcagt actgcggcgg gtagg 325

$$\begin{array}{ll} \langle 210 \rangle & 534 \\ \langle 211 \rangle & 334 \end{array}$$

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(334)
<223> n = A,T,C or G

<400> 534
ccgccagtgt gatggatatt tgcagaattc gcccttagcg agnnagccgg gcaggtccat 60
ggctaggttt atagatagtt ggggtggttg tggggnatga gtgaggcagg agtccgagga 120
ggttantttg tggcaataaa aatgattaag gatactagta taagagatca gggtcgtcct 180
ttagtggtgc gtatggctat catttggttt gagggtagnt tgattagnca ttgttgggng 240
gtaattantc ggctgttgat ganatatattg gaggtgggga tcaatanagg gggaaatana 300
atgatcagtn ctgcggcngg tnnacacctn gcc 334

<210> 535
<211> 557
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(557)
<223> n = A,T,C or G

<400> 535
nccataagct tcagtgcgca aaaggtcaag gccagtgtta atttgttatt tcttaaataa 60
ctttcccttt cattttttaa ttataaattt aacttctaac atgttttatg gttaaaattg 120
tacttttttc ctttagcgac attcaaattc atcacaatca ctttgtgaaa ttgttcgcct 180
gagcagagac cagatgttac aaattcagaa cagtacagag cccgaccccc tgcctgccac 240
tctagaaaag tatgtgtaaa actctgttct tgttcttctt tcatattgat gctgttccat 300
gtgttaccat tgtgagtggg tggttaagtgt tccttatgtg ggaatcatgt gccttgaaaa 360
taaccttggg tgggtgagaa ggtagggaaa cctgcttctt ttatctcaag taaaagtttt 420
ggcagggtaa agaagataaa tgacatttat atctagactt ttgagttttc caattatttg 480
gtaaaaatgg gaaattctgt agaagccctt ccttaaaaat gggggaagtc catttnanaa 540
aattaactgg taggtca 557

<210> 536
<211> 372
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(372)
<223> n = A,T,C or G

<400> 536
gttccaacct tcattttctga aactgttcta gagcacngtg tctttctcgt agttcataac 60
ttacccttct agtctagaat tagaattaca ttatctgttt tactacttta ctagactgta 120
agtccttaga agataaggac tagggagttc atctctgtat tccaccagaa ggtacagtga 180
ctcatatcta gagtcttttag atgaaactta ctgagttgaa taacttaata tatttctgtt 240
ttcattccca agggaggcca tgtctggaga tagacctga atttaataaa ttttaggcac 300

```
<210> 537
<211> 284
<212> DNA
<213> Homo sapien
```

<400> 537

```
<210> 538
<211> 293
<212> DNA
<213> Homo sapien
```

<400> 538

```
<210> 539
<211> 468
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(468)
<223> n = A,T,C or G
```

<400> 539

| | |
|-----------------------|-----|
| $\langle 210 \rangle$ | 540 |
| $\langle 211 \rangle$ | 397 |

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(397)
<223> n = A,T,C or G

```
<400> 540
ctgttttatt aattccccca tttgcagcac acttntctct tccaacattc atcagtcaga    60
tcagagtcca cggctcttttc aaaattttaga taaactggct tacattttgt aatgatgtcc    120
ccagacaaca cccactoca acccattctg tttgttacta ttagtttaca acatgcatgt    180
gcccttactt tcatttttcat agtattttaa aatggaaggg cactcccaaa tttactttaa    240
cccctttaat aatctctctc ctccctgctct ctctggctct ccagacaact gttgatttac    300
tttcctttat gatggattag tttgcatttt ctagaatttt atatgactga catataaagn    360
ttttatgttt ctcccctttg ggtttcttca tgtggca                                397
```

<210> 541
<211> 248
<212> DNA
<213> Homo sapien

```
<400> 541
cctagatagg ggattgtgcg gtgtgtgatg ctagggtaga atccgagtat gttggagaaa    60
taaaatgtgc atagtggggg ttttatttta agtttggttg ttaggtagtt gaggtctagg    120
gctgttagaa gtccctaggaa agtgacagcg agggctgtga gttttagggt gagggggatt    180
gttggttgga aggggggatgc gggggaaatg ttgtagcaaa tgagaaatcc tgcgaatagg    240
cttcgcggc                                248
```

<210> 542
<211> 366
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(366)
<223> n = A,T,C or G

```
<400> 542
aatcgccct ctagatgcat gctcgagcgg ccgccagtgt gatggatatt tgcagaattc    60
gcccttgagc gatanccggg gcaggtccaa ttgatttgat ggtaagggag ggatcgttga    120
ccnctctgt tatgtaaagg atgcgtaggg atgggagggc gatgaggact aggatgatgg    180
cgggcaggat agttcagacg gtttctatct cctgagcgtc tgagatgtta gtattagtta    240
gttttggtgt gagtgttagg aaaagggcat acaggactag gaagcagata aggaaaatga    300
ctatgagggc gtgatcatga aaggtgataa gctcttctat gataggggaa gtagcgtctt    360
gtanac                                366
```

<210> 543
<211> 460
<212> DNA
<213> Homo sapien

<400> 543

<221> misc_feature
 <222> (1)...(631)
 <223> n = A,T,C or G

<400> 553
 ccgggattag aactaaaaca agtgagatca cccctctaata ttttctgaa cttgggttaat 60
 aaaagtttat aagattttta tgaagcagcc actgtatgat attttaagca aatatgttat 120
 ttaaaatatt gatccttccc ttggaccacc ttcattgttag ttgggtatta taaataagag 180
 atacaaccat gaatatatta tgtttatata aaatcaatct gaacacaatt cataaagatt 240
 tctcttttat accttcctca ctggccccct ccacctgccc atagtcacca aattctgttt 300
 taaatcaatg acctaaagatc aacaatgaag tttttataa atgtatttat gctgctagac 360
 tgtgggtcaa atgtttccat ttcaaatata ttanaattc ttatgagttt aaaatttgta 420
 aatttctaaa tccaatcatg taaaatgaaa ctggtgctcc attggagtag tctccacct 480
 aaatatcaag atggctatat gctaaaaaga gaaaatatgg tcaagtctaa aatggctaata 540
 tgtcctatga tgctattatc atagactaac gacntttatc ttcaaaacac caaattgtct 600
 ttagaaaaat taatgtgatt acaggtagag g 631

<210> 554
 <211> 558
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(558)
 <223> n = A,T,C or G

<400> 554
 ccaggntagt ctccaactcc tgaccttagc tgatccaccc acctcggcct cccaaagtgc 60
 tgggattaca ggcattgagc actgcgcccg gccaaacttg atatgcattt ttaaataagt 120
 taatacatta ttcatggttt agtctcatta tatattctat ggtccacttt gaaatttcat 180
 ctaacaaaaa tcatcttcat cctgcaattt gaggtttgga cacaatgggg attgatcagt 240
 aatttcttca tatgcccttt ctcaaggaaa tagtttccta tgaaaaaaa gtcttatgtt 300
 ttcatgtaag ttctcttttt ggagaagaaa aggagacatt cttacttagc actctcagtt 360
 ttacaaaacg ctgccaacct taaaatttgt ctattgattc ccaaggcaca caaccaatag 420
 tctgtcaata acccggaata acatttcttt aaggccccag taactttcac atgtttgggt 480
 tccaatcttc acctagaatc ttgttaagaa aagtaaacca ttcactcttc tagaaactct 540
 aaggttgctt cttagggg 558

<210> 555
 <211> 212
 <212> DNA
 <213> Homo sapien

<400> 555
 ccaggatatt gcataatggc ttttcttctg ttgcctttgt tcctttgtgg cccagctaa 60
 ttgcctgaga gtgcactgt tagttttcaa ctctttctga tagaaaccct gtgtactaac 120
 atggaaatct taggtaatct gctttttcaa agcacaatgc agaatttatt ggcgggtggtg 180
 taactttaag aatatccgag aagccaccaa gg 212

<210> 556
 <211> 219
 <212> DNA
 <213> Homo sapien

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(488)
<223> n = A,T,C or G

<400> 559
ccccactgta ctccagcctg ggtgacccca tctcaaagaa gaaaagttac cagatgtcat 60
gggtaaagggt tgggtcttcaa gtggcctcat aagttgtctt gcattttaat tcaggaatt 120
cattggacca ataggttaca ttttcgttcc tttttgttt tggttcatct gttaagcagt 180
gggggcctaa ttactgctcc tttgtaaaaa cacatthttcc caaagaacac tgaattaccg 240
ttcaaaactgg ttgttgatgg gtaacaagggt ctgtttttgc tgccccaaaa gggcttaaca 300
atttaggcgg atagtttact taaaaaaaaa aatcctttgg agacatactg aaaatgcaaa 360
ctagtttcta aattatcaat tccctacatg aanaagcagt ttgccanagt ttagtctcan 420
aaaatgactg gttggctcta tttaaatcan aaccaattt ctacgcacct gccgcgccgg 480
ccaagggc 488

<210> 560
<211> 602
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(602)
<223> n = A,T,C or G

<400> 560
cctanttaag aattccttgc cttagtgggt aacaaggact aaacacagac aatgggtgaa 60
acacagacgc taattcacat aacagagagt aggcaacctt aagaatgaat tgatgcagac 120
tcctatagaa ttctctgtgt atgactgggt tcttattttc tcctccttgt atgtagtga 180
aatttcatca ttatgaatag ttcttggat ctttttttaa agttgtgaat gcgagtgttt 240
ggctttgtaa tacaactttt tagtatccag aagataacca gtgctctacc aataaagatc 300
ttttgatata aagggtttta acttctgcca gttcttactc atttttttca ggttttttat 360
acatttctta aacaacacat acattatgta aaatataaga attaatgtac attctcaagg 420
ccagattcag tgacaaaatg cactaccgca atctagtaac acatttactc cttgctgcat 480
ataagtggcg tgtaagaaat acagggtata ttgttttgtg atccatgcag taaatgttca 540
caaatatcag gcaacaact agacgntctt cagctactaa aattaactgt cccagtcaca 600
aa 602

<210> 561
<211> 683
<212> DNA
<213> Homo sapien

<400> 561
gtctatthttt aaaaagaaag aaaaaaacca cttttttata gtccttagct ttgccatatg 60
cccgccttaa gtggaaggaa agttaatcac ttaactatgt tttataaaaa gaaaaaagggt 120
cttggaatgc tattactgtt cacacaaagt atgattctgt ttgaataagg caaatgctcc 180
ttttttttaa aaaagacatt actgtaatat caaaaaccgt ggcagtttgt atacaactct 240
gggcttgatt tttttttaa aaacagaatg aattgatgtc ttattttata aatgtttctat 300
atttattagg agaaaacttt atattgcctt ttttatcaat catgtaacag gcttatagct 360

```

ttccaacaga gctgcttgcc aaacaat ttttgtttat taaacagtgc tgaaacaaac 420
aggatcagca tttacttaag atgttaagaa tgaggacttt taatcagccg aaccaagata 480
ttgttacctg tatgcattcc caaagtctag atgctcagta tgttcagtca tatctttcag 540
aatcagtga ccgattaccc tttttttggg attcactcta catctgcca cctagttcac 600
cttggttttg tgtctgctgt agaaggggaa cataacttgg ttaaaccgta gggattatca 660
ttgtatacat gctgtgaaca tgt 683

```

```

<210> 562
<211> 420
<212> DNA
<213> Homo sapien

```

```

<400> 562
gcactttttt tccagtaagg attcatctct tgcctccta tatggtcatt atattttata 60
ttttacatat ttataaacat gacatatgta tttatgttcc acaaagggct ttgaatagaa 120
tttacacata gagttccctg gggtgatgtg tttatcaaaa tggaagataa agtgaattaa 180
ttacttaaat atttaacact attgaataga aataatttcc ccaatattgc ttcattgattt 240
agacagtcta ttaaatgttt aagcaaggca ctagactaag tttattaaga caaattttgg 300
aatatgtgca gaaatatgac ctgggctaata gtacagagtc aaagctgggt gaatgggtgtt 360
atatagtgga ttcagattga tgtggcagtg gtgggttacac taggggcact aagggttatcc 420

```

```

<210> 563
<211> 482
<212> DNA
<213> Homo sapien

```

```

<400> 563
ctccacctta ctaccagaca accttagcca aaccattttac ccaaataaag tataggcgat 60
agaaattgaa acctggcgca atagatatag taccgcaagg gaaagatgaa aaattataac 120
caagcataat atagcaagga ctaaccctta taccttctgc ataataaatt aactagaaat 180
aactttgcaa ggagagccaa agctaagacc cccgaaacca gacgagctac ctaagaacag 240
ctaaaagagc acaccctct atgtagcaaa atagtgggaa gattttatagg tagagggcag 300
aaacctaccg ggcttggtga tagctgggtg tccaagatag aatcttagtt caactttaac 360
tttgcccaca gaaccctcta aatccccttg taaatttaac tgtagtcca aagaggaaca 420
gctctttgga cactaggaaa aaaccttgta gagagagtaa aaaatttaac acccatagta 480
gg 482

```

```

<210> 564
<211> 302
<212> DNA
<213> Homo sapien

```

```

<400> 564
ctggaagtga aggtactaat atacaaatgg ctcttgtttc tgaatatgtg atataatttg 60
tgaatctttg gaaactgaat tttttctatg gagtgcaaat atagaagggg tattttacaa 120
tggttggtgt gaaaagaatt cactttgtaa acaactatta aggctggaag tttagtgaag 180
gtgcatagtt ttgaaagcta cacagggtgaa aaatcaaact tattgtttgt aattttgctg 240
ttacatgtta agttactttg acagcaattt tctaatagata atgtgattta tgatttaaaa 300
gg 302

```

```

<210> 565
<211> 554
<212> DNA
<213> Homo sapien

```


<220>
 <221> misc_feature
 <222> (1)...(554)
 <223> n = A,T,C or G

<400> 565
 ccanngtgac atcatggcaa tacagcaaga attctgnnat ttatttagaa gcctcaagga 60
 gaaggatcct ggagcccctg aatgagagtt tcttctccat gcctctcccc agtcaaaaata 120
 catggaaaata ttcataagaag cattgtaccc agcatgataa ggaaggatgg agaattggtc 180
 cttatatctc tgttcacaag acatcaacac tcttaagtaa ctgtatgaaa taaattctct 240
 gctgaaagca aataaaccat ctgaaaggtc ttctgggttac ttacacagat ttcctagaga 300
 atctgaaatc agcctaagag ggaagattaa tttttaaatg aatccaagtt aatgaaagca 360
 aagaactctt atacagaaat acattttcct attataaagc aggactacct tccctaattt 420
 ctgatatgacc taggacaatt tgaatgggca ttgaaattct tttgggttgaa ttacgcaaac 480
 aagcaaagga aaagtctcaa ttattattgg aaaatttggg gagagattat tatctcttga 540
 tctcctagtn natt 554

<210> 566
 <211> 631
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(631)
 <223> n = A,T,C or G

<400> 566
 ncgaagctgt gaanncattc acacggaatc tgganggtat tactgtaact tcttataata 60
 cataatataa aagtttttga aagatataga cacaattaac ccctaaacaa cacactatct 120
 gattctcaaa agcaatggct atttaacaag atgtaaaagg acaataacat atcaaagaac 180
 tttcacacac ctaaagatag catttagcag caagttagtc agacaaaaca aacataaata 240
 tottcacatt tcttatgttt gtttttaact ttacttcata aagccactga taattgaggt 300
 ttcttttcaag tataagattt ctaaaattaa aaactgtttt tgacatatatt ttataaagaa 360
 ataaaaagca aaacgcaatc caactattta tatgagtcct tcttctccaa cagctttaga 420
 tgtttttctg agtacttttt acacagaata tttttattaa aatcagttct aattcattta 480
 tgcagattag gggaaaatga ttcataataa attaacctta aaattacctt ctatctgctt 540
 ctacctctat ccccccata ccaccaaatc tgttgctaca gtgaaactga gccaatgtct 600
 gtttgagggg gcccaaagca tctggtaatc t 631

<210> 567
 <211> 510
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(510)
 <223> n = A,T,C or G

<400> 567
 cctatnatag cttctctagc tatcatactc caatcagcna aaaatgagaa aatgttgaga 60
 aatagaagat aattcctcat ttaaggncac cttctanaat ttgtgcttaa nantctgttt 120

```

tcttctcatg ggccagcact tcggcaactg ggaaaaatta nngtacagg gatctaggna 180
atactgttta tttagagcaat aatatattgn gctaacgttc aggcataccta ttactgagaa 240
ataagggaaa atgagtgtaa agtacaacta agagtctcgg ctacaggga aaataccatc 300
agttaaatat ccatagtcct agagcattta tgtaaaactg caatttgaat cctgcaatac 360
attttggctt ttctctcagt gataccatgt gtgggaagtt gttctgtcaa ggtgggtcgg 420
ataatttgcc ctggaaagga cggatagtg ctttcttgac atgtaaaaca tttgatcctg 480
aagacacaag tcaagaaata ggcattggtg 510

```

```

<210> 568
<211> 180
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(180)
<223> n = A,T,C or G

```

```

<400> 568
ttaatntgac ncacgcttat gcggaggaga atgntttcat gttacttata ctaacattag 60
ttcttctata gggatgata ttgggtccaat tgggtgtgag gagttcagtt atatgtttgg 120
gatttttttag gtatgggtg ttgagcttga acgctttctt aattgggtggc tgcttttagg 180

```

```

<210> 569
<211> 237
<212> DNA
<213> Homo sapien

```

```

<400> 569
ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt 60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
atttctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt caggaaaagg 180
gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaag 237

```

```

<210> 570
<211> 352
<212> DNA
<213> Homo sapien

```

```

<400> 570
ctgtctctcc atttagagcc ccagttggtc ctgacctctt acaaatttgg tgttttccact 60
ttgatgttta tgaaccgatt gcattaaaaa tgcaggataa tgattcaggg ttagagaaac 120
tattatttat acaaattgtg ttaacacctc atcattttta attggctgtg ctaataatgc 180
tcattgtgct cttcaggggt atgtgtgtgt gtgtgtgtgt gttttgcctg aatctgcaac 240
ctacatttgc tctggcagta tgttgagtat atgctagaat agaattggacc taggcaactc 300
taaggctcta caactaaata cacttactta ggaaacctcc taaataagta gg 352

```

```

<210> 571
<211> 402
<212> DNA
<213> Homo sapien

```

```

<400> 571
ctgattttta caataactac tgtgttctct gcaatagtgt gttctgatta gaaatgacca 60

```

```

atattatact aagaaaagat acgactttat tttctggtag atagaaataa atagctatat 120
ccatgtactg tagtttttct tcaacatcaa tgttcattgt aatgttactg atcatgcatt 180
gttgagggtg tctgaatggt ctgacattaa cagttttcca tgaaaacggt ttattgtgtt 240
tttaatttat ttattaagat ggattctcag atatttatat ttttatttta tttgtttcta 300
ccttgagggtc ttttgacatg tggaaagtga atttgaatga aaaatttaag cattgtttgc 360
ttattgttcc aagacattgt caataaaagc atttaagttg aa 402

```

```

<210> 572
<211> 70
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(70)
<223> n = A,T,C or G

```

```

<400> 572
tggatccgag ctcggtacca agcttggcgt aatcatgggc atagctgttt cctgtgntcg 60
ttttacaacg 70

```

```

<210> 573
<211> 423
<212> DNA
<213> Homo sapien

```

```

<400> 573
ccaatgggtt cttagtgaag gagtacctta gctctgaatg caatgccttc agaaagatat 60
cattcataga gacatacaaa gcacatggca acatgacatt ggaatacacg attctgagca 120
tcttcattca tgaccaacct ggctatagat ttcagatgtc ctcttgggtc gaaggatata 180
tgggatatcc atgctcactt gcattccttt ccctttaatt tcattttcta agtccttctt 240
gtattgtttc taaaagaaca gaaaataatc ttggagcttt gcttaagctt taatagcgat 300
gttgaaattt acatgtttga atctcaaagc caccatgtg gaaagaaaac ttatgctctt 360
tccagctatg attcacggca tttattttta actttgtatc ttgctgctgt cttacctggc 420
tgg 423

```

```

<210> 574
<211> 129
<212> DNA
<213> Homo sapien

```

```

<400> 574
ctgttaaaag aacaaactta gcaatatata acagtttgct aacaggattt ttgactattc 60
actttgcgag ttatttttta aaatccactt ttttactgag ttttactaca taccaggcac 120
tgtacttgg 129

```

```

<210> 575
<211> 684
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(684)

```

<223> n = A,T,C or G

<400> 575

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccagatntga | cttttcaaaa | ctactcacat | tgtgaaaaan | gcaggaacaa | atctagtttc | 60 |
| aagttcagca | tgccgttccc | tgtttaattc | ataaaacaca | actggcagaa | gtattacttg | 120 |
| aagcaaaaca | aaagtaacgt | gggaacttgc | ttatttgcta | agccacaatg | tatttttcca | 180 |
| ggaatagcat | aaatttgcca | tctttcttgt | gtctatggaa | aaggggttta | gaattgtttc | 240 |
| actaaaaatt | aaatttctat | attgtcaaac | atgattgtat | actcaaattt | taaaatgtga | 300 |
| agggaacact | tactaagcat | ttcctgggta | tgccactata | ttaagtccta | gtaatatgat | 360 |
| atagtttatt | tcaatttttt | ttcaactcat | acttccttta | aaatagcact | gacccaaaaga | 420 |
| aagttaacat | gagcttcatg | tacaattttt | aatctttttg | cagaaaaata | aactgagaaa | 480 |
| ggctaaaatt | gttttattta | agccactata | ccaagacata | ttgatttcac | caatataaaa | 540 |
| attgagatag | tttacatttt | ttgggtacac | tttaaaatct | ggatgtattt | tttatactga | 600 |
| cagcacatct | caatttggac | aagctacatt | tccagggctc | aatagtcacc | atgaatctca | 660 |
| attgtaatca | aagaggttgg | cctg | | | | 684 |

<210> 576

<211> 134

<212> DNA

<213> Homo sapien

<400> 576

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccttatttct | cttgtccttt | cgtacagggg | ggaatttgaa | gtagatagaa | accgacctgg | 60 |
| attactccgg | tctgaactca | gatcacgtag | gactttaatc | gttgaacaaa | cgaaccttta | 120 |
| atagcggctg | cacc | | | | | 134 |

<210> 577

<211> 133

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(133)

<223> n = A,T,C or G

<400> 577

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtctctcc | attnagaagc | cccantnggt | cctnacctct | tacaaatttg | gtgttttcac | 60 |
| tttgatgttt | atgaaccgat | tgcatataaa | atgcaggata | atgattcagg | gttaganaaa | 120 |
| ctattattta | tac | | | | | 133 |

<210> 578

<211> 200

<212> DNA

<213> Homo sapien

<400> 578

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctcaaattc | atcttcaaa | gtgaccagc | aatcagtgtc | aatgccttta | ctgtagttaa | 60 |
| cctggtaatt | tcattcttta | gtctctccaa | gaaaatctga | agtgtattag | gcaagtcaga | 120 |
| acccaaattg | tctccaaggt | tgcaataaat | ttgtcccata | caggaaatag | ccctttcctt | 180 |
| gacttcctga | tcaatgtcag | | | | | 200 |

<210> 579

<211> 402

<213> Homo sapien

| | | | | | | |
|---|---|---|---|---|---|-----|
| c | c | c | a | a | a | 60 |
| g | t | t | a | a | a | 120 |
| t | a | a | a | a | a | 180 |
| t | t | a | a | a | a | 240 |
| a | a | a | a | a | a | 300 |
| t | t | a | a | a | a | 360 |
| g | a | a | a | a | a | 420 |
| c | a | a | a | a | a | 480 |
| g | | | | | | 481 |

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccaattaaga | gctaaattta | caaaataatc | tctatcagga | ggctttaagg | tttaatgtct | 60 |
| ctaaagtcct | tatggatata | agaggcttga | atgtactgaa | ttcaaatttg | gtttttaaat | 120 |
| gttataatag | tttaggcccc | agagccacat | atttctgtct | aagaatagaa | agcatagcta | 180 |
| gctgccacac | cagaatattc | atatagaggt | ggggggcaag | aacaaaattt | attcatttga | 240 |
| tacatagaaa | tgggactact | tagaatagac | tcataataga | aagcatcatc | tggttttctca | 300 |
| tctcag | | | | | | 306 |

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| ccagaatggt | acagagtgga | gggtgttctg | ctaattgactt | cagagaagta | tttaagaaaa | 60 |
| acatagaaaa | acgtgtgcgg | agtttgccag | aaatagatgg | cttgagcaaa | gagacgggtgt | 120 |
| tgagctcatg | gatagccaaa | tatgatgcca | tttacagagg | tgaagaggac | ttgtgcaaac | 180 |
| agccaaatag | aatggcccta | agtgcagtgt | ctgaacttat | tctgagcaag | gaacaactct | 240 |
| atgaaatgtt | tcagcagatt | ctgggtatta | aaaaactaga | acaccagctc | ctttataatg | 300 |
| catgtcag | | | | | | 308 |

<213> Homo sapien

<223> n = A, T, C or G

| | | | | | | |
|------------|------------|-------------|------------|------------|-------------|-----|
| cctgtctttg | aatggatgaa | atagggttaat | aaaaaacatc | actgtttaaa | aactagaaca | 60 |
| ctgaaaaatt | ctaggaaagc | ttattttccc | ttatatTTTT | atggnaCTTT | caacacttna | 120 |
| caacactatt | tnaattaann | tttnttctag | agtttatann | atatcagtac | attctttttct | 180 |

gtggatgcaa taatatagaa tcttattnca aatcttactg gcaggntctn ttaaattctt 240
 caacggntgn catagtgatt aacccaaaatt agttatgatt tctgcctatc tgtgtgagaa 300
 cttacagggg aaattgttct aaacctgagg aacatgaagt aactgtactg cacactccaa 360
 atgatgacag tcattttata tcaccttcaa ttaccaaca gcttttaata gtctgg 416

<210> 587
 <211> 382
 <212> DNA
 <213> Homo sapien

<400> 587
 cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga 60
 gctgttcctc ttgggactaa cagttaaatt tacaagggga tttagagggg tctgtgggca 120
 aattttaaagt tgaactaaga ttctatcttg gacaaccagc tatcaccagg ctcggtaggt 180
 ttgtcgctc tacctataaa tcttcccact attttgetac atagaagggt gtgctctttt 240
 agctgttctt aggtagctcg tctggtttcg ggggtcttag ctttggctct ccttgcaaag 300
 ttatttctag ttaattcatt atgcagaagg tataggggtt agtccttgct atattatgct 360
 tggttataat ttttcatctt tc 382

<210> 588
 <211> 307
 <212> DNA
 <213> Homo sapien

<400> 588
 cctactcttc tccgtccatt gtactatctg cccgtgggtg ggatggcagt aggatcatat 60
 ttgatgactt ccgagaagca tattattggc ttctgcataa tactccagag gatgcgaagg 120
 tcatgtcctg gtgggattat ggctatcaga ttacagctat ggcaaaccga acaattttag 180
 tggacaataa cacatggact aataccata tttctcgagt agggcaggca atggcgcca 240
 cagaggaaaa agcctatgag atcatgaggg agctcgatgt cagctatgtg ctggtcattt 300
 ttggagg 307

<210> 589
 <211> 89
 <212> DNA
 <213> Homo sapien

<400> 589
 cctgggtgat tgaggatgca atgagctgtg attgtgccac cacactccag cctgggcaat 60
 acagcaagac tgtctcaaaa aaaaaaaaa 89

<210> 590
 <211> 456
 <212> DNA
 <213> Homo sapien

<400> 590
 cctcagttct tgatttgtgt tgacggggcg tcaccatgaa ggagcccatt tagtataaag 60
 cttccaacct tttctcttaa tegtctctt aatcttttaa accatcttca agtgcatagg 120
 ggagtttccg atgccagagg atgaaagcaa gtgctctctc caccctctcc tcccagagt 180
 aaaacaaatc cttttgtctga tacttgtttc aaaagcatcc attgtaaagc ttctcagtga 240
 cacaaaatac tgagaggtaa ctttttatca atcaaaccac ataccccaat ttaacacctt 300
 tcaatgctct gaattcaact gacagactaa aggggtgttc ctgtaacagt ctgaaatatt 360
 aagtgttttt tttgttttgt ttttaaactc tatttcagaa aacttctct tggggtagga 420

aagtacacat gaagcagcaa agtaacgaag aaaaac

456

<210> 591
 <211> 289
 <212> DNA
 <213> Homo sapien

<400> 591
 ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt 60
 agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
 atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
 gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaagggtg 240
 ataagctctt ctatgatagg ggaagtagcg tcttgtagac ctacttgcg 289

<210> 592
 <211> 435
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(435)
 <223> n = A,T,C or G

<400> 592
 cgcgttagat gcgccttttc cggcctgtgc gtctgctctg gttcctctca ggcagcaaag 60
 ctggggaagg aagctcaggc aggagcctcc ccgacaccac agcggcacia gcagcagcta 120
 aagcaccgca ctttgctctg ctaacctttt acttaaatga ggttttgcca aatccacatc 180
 tggaaccgca tcacacccat ttgcaaggat gtttggtctt tgatgaaact gcatctctac 240
 tgcacatgan ggctttcatt gtaggacaag aggagagtgc gtttatTTTT gtaactgttt 300
 tacatgttcc gattanttaa tcggnagctt atgtcatttg ctatgcctgt tgtcttctaa 360
 tctctcctta ctaaaacatt acttcaaatt tnaattgacc cttgtttata atttatttaa 420
 cgggatttgn gtgtc 435

<210> 593
 <211> 633
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(633)
 <223> n = A,T,C or G

<400> 593
 ctgttttagtc agataattgt gtccgaattg attangaaaa taatagacca gccataaagc 60
 agcataaaaat attatgaaac tattccagaa gttcagtaat atcttttgga cctgtctata 120
 gcccaagttt tgtgaatact tttgtagtta aaaaaattt ttactttacc agggcattgc 180
 aattcttttc catcagtga tttcattcta cagacttttc agagcatctc ataatacgtc 240
 aacaaatcta tttcaaagt gtttgttact aagcaacggt tgctaagagc ttctgtaatt 300
 aagatgaaag ttccaaggta acaatgccca aacacagcac cattttcacc attttctgat 360
 aatgcaggag taggatggct aaaagtgaag gaagaatcta ctctatggaa agcatggcac 420
 ctgaaatttc tgaagatatt ggctgtcctc tagcttatat gagagagagt gtttgtgctt 480
 tactaatcaa ccagtcattt ttttcttggt tggctgaaat gtacattcca gacatgaaca 540

ggtagagtat gtgttggggg caggtttata ctgcatgggt gtgctgagac agggccacgt 600
 ggtgatgtaa atgatgctgn ctgacacgtg cag 633

<210> 594
 <211> 501
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(501)
 <223> n = A,T,C or G

<400> 594
 cctttacaag atgctggtac cttgatcttg gaconggcag gctccaagat ggaaagaaag 60
 tgagcatctg ctttttaggg attatccagt ctatactact ctgttctagc cacacaaaac 120
 aggttaagac agaaattggt accaagagtg ggggtgttact acagcaaata cctgaaaatg 180
 tagaagaggc tttgaaatgt ggtaattgga agaagctggt agaatttgga ggagtaggct 240
 agaaaatgtc tgtattttca tgaatggagc attaagaata attccggtga ggccataggg 300
 aaagtctaaa actttttcaga aattatgtaa gcgattgtga ttagtagggtt ggtagaaata 360
 tagacagtaa aagcaattct gatgtggttt cagaggaaaa tgaaaaatat tagaaactga 420
 aggaaggggc atccttgcta taaactggca aagaacttgg ctgaaatgtc tccatgtcca 480
 agagatttat ggcagaaatg t 501

<210> 595
 <211> 383
 <212> DNA
 <213> Homo sapien

<400> 595
 ctggtcacca tcatcccttt aatcaactca cacctgttta aagagtgttt ctgatttgac 60
 cttcatccct tagtttactg gcgttaaaaa aagtctcagc aattttcatt atttctcgtg 120
 ggtctcatta tcaaacccttt acttatttcg gcataatttc tctgggcttc ttctagtttc 180
 tgccttacia gcaatgctgt tctgtaaatt tattgaaacc tctggaacat ttcaccttta 240
 gagatggagg atggaaggat tgggtaccaga agagggtctaa gatacgtttt ctgtcttgag 300
 ctgaaagcac agtctactct ccttcgtttt gtcgatgaga aagttgaggc cagagggggag 360
 gtgacatgtt tagagtcacc cag 383

<210> 596
 <211> 266
 <212> DNA
 <213> Homo sapien

<400> 596
 ccattggctag gtttatagat agttgggtgg ttggggtaaa tgagtgaggc aggagtccga 60
 ggaggttagt tgtggcaata aaaatgatta aggatactag tataagagat caggttcgtc 120
 ctttagtggt gtgtatggct atcatttggt ttgaggttag tttgattagt cattgttggg 180
 tggtaattag tcggttggtg atgagatatt tggaggtggg gatcaataga gggggaaata 240
 gaatgatcag tactgcggcg ggtagg 266

<210> 597
 <211> 383
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(383)
 <223> n = A,T,C or G

<400> 597
 ctggtcacca tcatcccttt aatcaactca caccngttta aagagtgttt ctgatttgac 60
 cttcatccct tagtttactg gcgttaaaaa aagtctcagc aattttcatt atttctcgtg 120
 ggtctcatta tcaaacccttt acttatttctg gcataatttc tctgggcttc ttctagtttc 180
 tgccttaca gcaatgctgt tctgtaaatt tattgaaacc tctggaacat ttcaccttta 240
 gagatggagg atggaaggat tgggtaccaga agaggggctaa gatacgtttt ctgtcttgag 300
 ctgaaagcac agtctactct ccttcgtttt gtcgatgaga aagttgaggc cagaggggag 360
 gtgacatggt tagagtcacc cag 383

<210> 598
 <211> 266
 <212> DNA
 <213> Homo sapien

<400> 598
 ccatggctag gtttatagat agttgggtgg ttggtgtaaa tgagtgaggc aggagtccga 60
 ggagggttagt tgtggcaata aaaatgatta aggatactag tataagagat caggttcgtc 120
 ctttagtggt gtgtatggct atcatttgtt ttgagggttag tttgattagt cattgttggg 180
 tggttaattag tcggttggtg atgagatatt tggaggtggg gatcaataga gggggaaata 240
 gaatgatcag tactgcggcg ggtagg 266

<210> 599
 <211> 294
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(294)
 <223> n = A,T,C or G

<400> 599
 ccaattgatt tgatggtaag ggagggatcg ttgaccacgt ctgttatgta aaggatgcgt 60
 agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
 atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
 gcatacagga ctaggaagca nataaggaaa atgactatga gggcgtgatc atgaaagggtg 240
 ataagctctt ctatgatagg ggaagtagcg tcttgtagac ctacttgccg tgca 294

<210> 600
 <211> 213
 <212> DNA
 <213> Homo sapien

<400> 600
 agatattggg ctgttaattg tcagttcagt gttttaatct gacgcaggct tatgcggagg 60
 agaatgtttt catgttactt atactaacat tagttcttct atagggtgat agattgggtcc 120
 aattgggtgt gaggagttca gttatatgtt tgggattttt taggtagtgg gtgttgagct 180
 tgaacgcttt cttaattggt ggctgccttt agg 213

<210> 601
 <211> 471
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(471)
 <223> n = A,T,C or G

<400> 601
 ncctactatg ggtgttaaatt tttttactct ctctacaagg ttttttccta gtgtccaaag 60
 agctgttcct ctttggaacta acagttaaat ttacaagggg atttagaggg ttctgtgggc 120
 aaatttaaag ttgaactaag attctatctt ggacaaccag ctatcaccag gctcggtagg 180
 tttgtgcct ctacctataa atcttccac tttttgcta catagacggg tgtgctcttt 240
 tagctgttct taggtagctc gtctggttc gggggtctta gctttggctc tccttgcaaa 300
 gttatttcta gttaattcat tatgcagaag gtataggggt tagtccttgc tatattatgc 360
 ttgggtataa tttttcatct ttcccttgcg gtactatata tattgcgcca gggttcaatt 420
 tctatgcct atactttatt tgggtaaatt gtttggtctaa ggttgctctgg t 471

<210> 602
 <211> 482
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(482)
 <223> n = A,T,C or G

<400> 602
 tgagcataca gcaataaaaa taacataatt tntatgtgta caatatattat ggaatacgtt 60
 actggaacag ataaataatt tagttaataa catgacaaag aacagaaatt gtatacacta 120
 tacagcatag taatagaata atgaatgatt aaagttatta atattaggta gaaaatgaag 180
 ggtatctttg agagcagaac tcaaggaagc aagcaatttg ccttatgagg aaagagttac 240
 ctgtggataa aggagaaact gaaaaattta caagtcaaga ctttttgagc aaaaacaaaa 300
 atatgactat gagtcaccaa ttcagtacag tgaaaaaaa gttgaagaga tatcttggaa 360
 gtaaaccatg ttgtggaaga gcagggtttt gataatcatg ggattattct gaatgaattt 420
 taaatgcgat aggaatatat gagataattt caccagagaa taatatgatc atgtttgcat 480
 tt 482

<210> 603
 <211> 372
 <212> DNA
 <213> Homo sapien

<400> 603
 gttccaacct tcatttctga aactgttcta gagcactttg tctttctcgt agttcataac 60
 ttaccccttc agtctagaat tagaattaca ttatctgttt tactacttta ctgactgta 120
 agctcctaga agataaggac tagggagttc atctctgtat tccaccagaa ggtacagtga 180
 ctcataacta gagtcttttag atgaaactta ctgagttgaa taacttaata ttttctgtt 240
 ttcatccca agggaggcca tgtctggaga tagaccttga atttaataaa ttttaggcac 300
 tataccattt cagtggagaa aattgttggg aaatttgggg ggatggatat ataaggggga 360

ggaagtcact gg

372

<210> 604
 <211> 468
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(468)
 <223> n = A,T,C or G

<400> 604
 gcngttttga gtgagtttct taatcctgag ttctggnttg attgcactgt ggtctgagag 60
 atagttttgtt ataatttctg ttctttttaca cttactgagg agagctttac ttccaagtat 120
 gtgggtcgatt ttggaatagg tgtggtgtcg tgctgaaaag aatgtatatt ctggttgattt 180
 ggggtggaga gttctgtana tgtctattag gtccgcttggt tgcagagttg agttcaattc 240
 ctggatagcc ttgttaactt tctgtctcgt tgatctgtct aatggttgaca gtgggggtgg 300
 aaagtctccc attattattg tgtgggagtc taagtctctt tgtagggtcac taaggacttg 360
 ctttatgaat ctgggtgctc ctgcattggg tgcacatata tttaggacag cnagctcttc 420
 ttgttgaatt gatcccttta ccattatgta atggccttgn ctcttttg 468

<210> 605
 <211> 288
 <212> DNA
 <213> Homo sapien

<400> 605
 ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctggttatgta aaggatgcgt 60
 agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
 atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
 gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgac atgaaagggtg 240
 ataagctctt ctatgatagg ggaagtagcg tctttagtagac ctacttgc 288

<210> 606
 <211> 572
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(572)
 <223> n = A,T,C or G

<400> 606
 gaatnaaatg aatgaaatag aaaatataat tgagagcttc aacaacagac tataccaaat 60
 ggaggaaaaa atttctgaac ttgaagatag atcttttgaa ataacacaag cagtggcaaa 120
 aatgaattaa aaagaataag gaaagcctaa aggatttatg agatatcatt aagcaagcaa 180
 atattcatac tatgggcatt ccagatggaa aaaagaaggg taaagggtgag gaaatcatat 240
 ttaatgaaat aatagcagaa aatttcaggga gtcttgggag agagatgagc atttaggtcc 300
 agggagctca aagaacccca aacagattca acccaaacag gtcctctctg gagcccaaca 360
 tagtcaaatt gtaataagta aaagacaaag aattccaana agcattcaag agaaaagagt 420
 caagtcataa ataagggaat ctccattagg ctaacagcag atatctcagc agaaagctta 480
 cangccanga gagaatggga tgatatattc aaagtacttg aaagcagggg tnggggaaac 540

cctgctagct aaaaatatta tacccttgca aa

572

<210> 607
<211> 178
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(178)
<223> n = A,T,C or G

| | |
|---|-----|
| <400> 607 | |
| ctcggggtaa tctcccagca agaggtcagg tcctggntgt gcgtcccagg gtgtcagtga | 60 |
| aattggctgc tcccctgacc cagggcacct tcatgcgtct tcacagcagg actactgtga | 120 |
| ccaaggccag acctttcatc tttcaaaaga ctttgactaa aaatgcttta aaaaagca | 178 |

<210> 608
<211> 416
<212> DNA
<213> Homo sapien

| | |
|--|-----|
| <400> 608 | |
| cctgtctttg aatggatgaa atagggttaat aaagaacatc actgttttaa aactagaaca | 60 |
| ctgaaaaatt ctaggaaagc ttattttccc ttatatTTTT atggtaacttt caacacttaa | 120 |
| taacactatt tcaattaagt tttctcctag agtttatagt atatcagtag attcctttct | 180 |
| gtggatgcaa taatatagaa tcttattcca aatcttactg gcaggttctc tttaaattctt | 240 |
| caacggctgt catagtgatt aaccaaaatt agttatgatt tctgcctatc tgtgtgagaa | 300 |
| cttacagggg aaattgttct aaacctgagg aacatgaagt aactgtactg cacactccaa | 360 |
| atgatgacag tcattttata tcaccttcaa ttacccaaca gcttttaata gtctgg | 416 |

<210> 609
<211> 648
<212> DNA
<213> Homo sapien

| | |
|--|-----|
| <400> 609 | |
| ctgatctctc agcagaaact cttcaaacca gaagagagtg ggggcccaata ttcaacattc | 60 |
| ttaaagaaaa taattttcaa ccagaaattt catatccagc caaactaacc ttcacaagtg | 120 |
| aaggagaaat aaaatccttt acagacaagc aaatgctgag agattttatc accaccaggc | 180 |
| ctaccctaaa agagttcctg aaggaagcac taaacatgga aaggaacaac cagtaccatc | 240 |
| gaggctagga agaaaccgca tcaactaagg agcaaaataa ccagctaaca tcataatgac | 300 |
| aggatcagat tcacacataa cgatattaac tttaaatgta aatggactaa atgctccaat | 360 |
| taaaagacac agactggcaa attggataaa gagtcaagac ccatcagggt gctgtattca | 420 |
| ggaaacccat ctcaccgtgc agagacacac ataggctcaa aataaagggc tggaggaaga | 480 |
| tctaccaagc aaatggaaaa caaaaaaagg caggggttgc aatcctagtc tctgataaaa | 540 |
| cagactttta accaacaag atcagaagag acaagaagg ccattacata atggtaaaagg | 600 |
| gatcaattca acaagaagag ctaactatcc taaatatata ttgcaccc | 648 |

<210> 610
<211> 310
<212> DNA
<213> Homo sapien

<400> 610
 ccagctcttc tctgtcacat tcctatttct gacttctgcc tggctttcag tttctgcccc 60
 accttggtt tttcccagct tgaacctaat agaactccag agtttggggg gagggccagc 120
 cctttgtttt ctgctcttga agcatattca cacataaaaa gttgtattct cttacacaaa 180
 ctgttttgag gctcttaccg tagtogaagg tatcttagat cttccttagt gatctcatta 240
 agaatatccg aaagtgtata accctcttca acaatctgaa acaaagatca gatccttaag 300
 agctgagcag 310

<210> 611
 <211> 254
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(254)
 <223> n = A,T,C or G

<400> 611
 ctgtttttac atctaaagca atagactaga actgaattnt cttctacata gtaaaatcac 60
 aattgtggaa ttacaggaat tctgggtgata ttaaggtgaa acaacaaaac acaaaaggcc 120
 ctattttaac agttgatgtg acagtaagtt ttaatagaac ctgtaacttc attttggaag 180
 tgcttctcca ccaaataaag cctttttccc ctatttaagg agccagatgg attgaaagat 240
 gtggaaatag gcag 254

<210> 612
 <211> 225
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(225)
 <223> n = A,T,C or G

<400> 612
 ctgactatat catgtcacca tcatagccaa tacaacattn ttgccatact tcctaaaaac 60
 cttttcgcat acactgatca tgctacttat cagcactttc taacatcctg accaaacaga 120
 caccacacc tcttatagag tacactgtga gagaataaca tggacttgat atggcatcac 180
 acttgtttta aagcaaaaaa aaaagaaaaa gaaaagaaaa aaaaa 225

<210> 613
 <211> 471
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(471)
 <223> n = A,T,C or G

<400> 613
 ccacagact tcttggtgct ctggctatat tcaatgtgaa gtaaaaaata tcccaagtct 60
 tacaccaaaa tagaggctct gacttagaag tatgctttta gctttctttt taaataagac 120

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccaccaagc | cacacggaga | ttctgtcagg | cgctgagaca | ccacagcctt | ttcaatctta | 60 |
| gggaaagaaa | tcaagtcata | taaattaata | tcaacaggta | aggtcattga | gcaattgtct | 120 |
| ttcaactgtc | taagacttta | tcacttaaga | tcataaacac | agaagcaggt | cataaaaaata | 180 |
| gcttttctta | aggtttagga | gaatttgtag | gggcacttac | ttgataatct | gaattttcta | 240 |
| gtcagaagtt | taaataccac | cttttaaaaa | cataaaattt | aatttgtaac | aagttattaa | 300 |
| caaagcagta | ttgtcgaag | ttttaagctt | tctccaata | atttaattac | attaattaaa | 360 |

tttttaccat tctaattggtt acaaagtaac cag

393

<210> 621
<211> 563
<212> DNA
<213> Homo sapien

<400> 621

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgacaatga | taaaattatc | tctatatggg | caaacgcgtg | ctctttgtcg | aagaagaaag | 60 |
| cttcagcttc | atgttccagg | tgagttaatt | aggcaatgta | tgaatgctaa | tatctctttc | 120 |
| acataatttg | cttaagatct | gtcttaggac | tctcgtcttg | cccatatggt | tttccaaggg | 180 |
| cagaagggcc | tctttttgat | gagaggcagt | tttcagtaac | tcttaaagtg | ataacagcaa | 240 |
| aggagaggag | agagaagagt | aagacaaatc | gaaacattct | tcaattgctt | cttggccttt | 300 |
| tggctaagct | caagctcaaa | acaggtcttc | aaggagaaaa | tacatcacaa | agaaaaggat | 360 |
| gttttatttc | ttaccttgct | ctagaaaaat | ttccataaac | tctattggct | taattctgta | 420 |
| aacttgacca | atatcagagt | gcttcctacc | aaggagggtg | gctgatgagc | gtgaccatgg | 480 |
| tacatcctag | aagaatgtgt | gatgaagaag | ctttcacctg | gtaaaagagt | tgaaaattat | 540 |
| tcaaggagac | attatggtct | tgg | | | | 563 |

<210> 622
<211> 505
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(505)
<223> n = A,T,C or G

<400> 622

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| tcttaagtgt | gtttaataga | taaagtaaac | tttcctagtc | aagggttaga | tttttattat | 60 |
| ctcttggtgt | ccgactttct | acttttcaac | tttgaacttc | aaaaaaacat | tactttgctt | 120 |
| atcctttgta | ctttgatcag | gttggttaga | attgtagatc | aaaccattct | ttgatcattt | 180 |
| tattgtttta | atgnttagtt | ccatttataa | tttttatagc | caactctcgg | ttattttctgt | 240 |
| cttttgagat | tgcaattcag | aagctgtatg | tcgaagtaat | ttatgagttg | actttttatac | 300 |
| ttaggcttct | ttaaatacta | atagtcaaga | attctagagc | atctaataaa | aaattaactt | 360 |
| tcagatcatt | gggaatctgt | cctcatttaa | atatgtgtaa | atgcatttcc | acagcaaatt | 420 |
| gcttcatgcc | ctttgnctat | aaggaaatta | ttccttgtag | ctaatacatt | tttcattttg | 480 |
| cagnccaaat | cttttttgag | aaagg | | | | 505 |

<210> 623
<211> 489
<212> DNA
<213> Homo sapien

<400> 623

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| cctactatgg | gtgttaaatt | ttttactctc | tctacaaggt | tttttcctag | tgtccaaaga | 60 |
| gctgttcctc | tttgactaa | cagttaaatt | tacaagggga | tttagagggg | tctgtgggca | 120 |
| aattttaaagt | tgaactaaga | ttctatcttg | gacaaccagc | tatcaccagg | ctcggtaggt | 180 |
| ttgtgcctc | tacctataaa | tcttccact | atcttgctac | atagacgggt | gtgctctttt | 240 |
| agctgttctt | aggtagctcg | tctggtttcg | ggggtcttag | ctttggctct | ccttgcaaag | 300 |
| ttattttctag | ttaattcatt | atgcagaagg | tataggggtt | agtccttgct | atattatgct | 360 |
| tggttataat | ttttcatctt | tccttgccg | tactatatct | attgcgccag | gtttcaattt | 420 |
| ctatcgctat | actttatttg | ggtaaattgg | ttggctaagg | ttgtctggta | gtaagggtgga | 480 |

489

<400> 624

gggaac agct.

aatta ggct

cggtg gtaa

acgta gtctaggcca t

<210> 625

<211> 459

<212> DNA

<213> Homo sapien

<400> 625

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| ttcgagaaca | tttttaataa | ataatgtgac | aaaattactt | ttctgattat | tggattttca | 60 |
| gtatgcaaaa | ttatggctaa | aaataagggg | cttcttacat | gaacataatg | aaaacattaa | 120 |
| tcacatggat | tgttccctta | gtactgcacg | ccttttctat | ggaacttttt | caaattatct | 180 |
| aaatgaacaa | gtttggtttt | ggtgaacacc | agcctttttt | tttgtggttc | agttttgttt | 240 |
| ggctttgtct | tccactgggg | tcagacctga | tacttatcta | tctatgaata | aatgtacatt | 300 |
| tttttcttca | aatagcacca | attataaaaat | caatgatatt | cataaaatga | caaaaaagga | 360 |
| tcatagaaat | ctactagtca | gagggcatca | tttgtcaatt | gaaagcaagt | aatgcctcta | 420 |
| ttagaqattt | taaggaaaac | ttgtaggttt | cgacattgg | | | 459 |

<210> 626

<211> 458

<212> DNA

<213> Homo sapien

<400> 626

| | | | | | | |
|-------------|------------|------------|-------------|------------|------------|-----|
| cctgatgatt | gttttaaac | gtagaaagg | ttcagctaag | aactacagtc | cactctcagc | 60 |
| cctgtcatgt | actataggac | aagtcttcat | tcacaacaaa | tggatagcaa | caccaatctc | 120 |
| gtaacactgg | gaaaactgca | tacaatattt | agaagggaaca | ctaatacagc | agaatctgca | 180 |
| cacaacggag | tcaaatgatc | gaggccaaat | cctactacac | tttacgactt | tgagttggtc | 240 |
| actttttctga | accttagctt | ctccatcagt | gtaaaactga | tgtaaaataa | tataaagcta | 300 |
| tatgaaagct | gatgtgattt | acttgtgaaa | tagtatgtgc | aaaaggactt | tgtaaaatgt | 360 |
| aaagcactat | gctggttatt | gtgatatctg | agatattttt | aaagttgcaa | ttcaattcaa | 420 |
| caagcattca | tttagagtca | tgtgcaaggc | actgtgct | | | 458 |

<210> 627

<211> 393

<212> DNA

<213> Homo sapien

 $\langle 220 \rangle$

<221> misc feature

<222> (1) ... (393)

<223> n = A, T, C or G

<400> 627

```

ccatnngaac gcactcagga ggtgggttgt tctggatgca gaaaccagag atctagtttc      60
tatccacaca gacgggaatg aacagctctc tgtgatgcg tactcaatag atggtacctt      120
cctggctgta ggatctcatg acaactttat ttacctctat gtagtctctg aaaatggaag      180
aaaatatagc agatatggaa ggtgcactgg acattccagc tacatcacac accttgactg      240
gtccccagac aacaagtata taatgtctaa ctcgaggagac tatgaaatat tgtactggga      300
cattccaaat ggctgcaaac taatcaggaa tcgatcggat tgtaaggaca tttgattgga      360
ccgacatata cctgtgggct aggacttcca gga                                     393

```

```

<210> 628
<211> 233
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(233)
<223> n = A,T,C or G

```

```

<400> 628
ctggatttat aaaatagttg aatgacaaaa gaagnntggt ttgacagtaa aaaaaagaca      60
ttatggacaa aatatgcaaa atgtgcaaag aaaaaataaa tttgcattag aaaggtgggc      120
atttgatctc tgagccctgt gccatgtaac attgccatgt tctttcactg ttgtttgaat      180
gttgtagccc ancccttgac tctggactta aggcaagcta tgactggctt tgg                                     233

```

```

<210> 629
<211> 450
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G

```

```

<400> 629
ccnggacaat ntaggcagga gaaggaaata aagggtattc aattaggaaa agaggaagtc      60
aaattgtccc tgtttgaga tgacatgatt gtatatctag aaaaccccat tgccctagcc      120
caaaatctcc ttaagctgat aagcaactcc agcaaagtcg caggatacaa aatcaatgga      180
caciaatcac aaacattctt atacaccaat aacagacaaa cagaggccaa atcacgagtn      240
gaactctatt ccaattgctt tcaagaaaat taaaatacct agggatccaa cttacaaggg      300
acatgaagga cctcttcaag gagaaactac aaaccactgc tcaatgaaat aaaagaggat      360
acaaagaaat ggaagaacat tccatgctca ttggtagctt gatgggggatg gcattgaatc      420
tataaattac cttgggcagt atggacctca                                     450

```

```

<210> 630
<211> 486
<212> DNA
<213> Homo sapien

```

```

<400> 630
cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga      60
gotgttcttc tttggactaa cagttaaatt tacaagggga ttttagagggt tctgtgggca      120
aattttaaagt tgaactaaga ttctatcttg gacaaccagc tatcaccagg ctcggtaggt      180
ttgtcgcttc tacctataaa tcttcccact attttgctac atagacgggt gtgctctttt      240

```


<400> 634

<210> 635

<212> DNA

<213> Homo sapien

<400> 635

<210> 636

<211> 383

<212> DNA

<213> Homo sapien

<400> 636

| | | | | | | |
|------------|-------------|------------|-------------|------------|------------|-----|
| cctactatgg | gtgttaaatt | ttttactctc | tctacaaggt | tttttcctag | tgtccaaaga | 60 |
| gctgttcctc | tttggaactaa | cagttaaatt | tacaagggga | tttagagggg | tctgtgggca | 120 |
| aatttaaagt | tgaactaaga | ttctatcttg | gacaaccagc | tatcaccagg | ctcggtaggt | 180 |
| ttgtcgcttc | tacctataaa | tcttccact | at ttgtctac | atagacgggt | gtgctctttt | 240 |
| agctgttctt | aggtagctcg | tctggtttcg | ggggctcttag | ctttggctct | ccttgcaaag | 300 |
| ttatttctag | ttaattcatt | atgcagaagg | tataggggtt | agtccttgct | atattatgct | 360 |
| tqgttataat | ttttcatctt | tcc | | | | 383 |

<210> 637

<211> 537

<212> DNA

<213> Homo sapien

<220>

<221> misc feature

 $\langle 222 \rangle \quad (1) \dots (537)$

<223> n = A, T, C or G

<400> 637

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ttttaatcct | ggggtatata | ggcagnactt | taaattgcaa | agtcttcg | gcctattttc | 60 |
| ctctacattt | ttgtaattaa | ctctgggggc | ttacttggtt | tggcagtact | gaaatcaaag | 120 |
| gagctggttc | ttctttttct | ccaattattt | tcatatgaaa | gcacctacaa | ttagcctggt | 180 |
| agtcctattc | agatacatca | aatatcagtg | aatgctttac | tattegcaca | tttaagcatc | 240 |

```

tttgttttac ataaaattag agtatgaaaa ccagtgttca attttttacc ttgttgagct 300
tgtaaaatgc cagcaattta aaactaggac ttttccccc ataagccaag gaggtagaat 360
tactaataca aggggttaaag aaggtagatt ttgttttcaa tatttgggta atattagaaa 420
gattcttccc acagggaaga actagcaagt gtcccaattt tttccaaacg ttggggaggg 480
gaaaattcac tgtatcatga aaccctaagg gtttgngtgc acttcctgct ttttagg 537

```

```

<210> 638
<211> 445
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(445)
<223> n = A,T,C or G

```

```

<400> 638
ccagcagaac acagnagtga tttgggtccc tttgttcccc agtgggggtat ctatccttgt 60
gcagggcaca agcctacatg gtggctctgg tcatatcatt agaaaataga cagaaatggg 120
ctgcacacca gaatgaatga attgaattga aaggaggagg tgatgggtgga aaaaaaaca 180
agtcaattca tttagactgg tagaaccaga accactgtgt agtacatcca aacgggttaa 240
attccctgga agatgttaca taatcctatc atgggtgttta tttatggaaa tctattttaa 300
aaattttatg taatactgca cagtctgttt gcatgatgcc ttgtacgtag tagcaactca 360
gtaaatactt tttgaatgaa ctagtatagt attttaatta gctagtcttc gtgtactggg 420
acaaaagaac agtgtcatct tacag 445

```

```

<210> 639
<211> 584
<212> DNA
<213> Homo sapien

```

```

<400> 639
gcttgagtat tctatagtgt cacctaaata gcttggcgta atcatggtea tagctgtttc 60
ctgtgtgaaa ttgttatccg ctacacaattc cacacaacat acgagccgga agcataaagt 120
gtaaagcctg ggggtgcctaa tgagtgaagt aactcacatt aattgcgttg cgctcactgc 180
ccgctttcca gtcgggaaac ctgtcgtgcc agctgcatta atgaatcggc caacgcgcgg 240
ggagaggcgg tttgcgtatt gggcgctctt ccgcttcttc gctcactgac tcgctgcgct 300
cggtcgttcg gctgcggcga gcggtatcag ctactcaaaa ggcggtaata cggttatcca 360
cagaatcagg ggataacgca ggaaagaaca tgtgagcaaa aggccagcaa aaggccagga 420
accgtaaaaa ggccgcgttg ctggcggttt tccataggct ccgccccctt gacgagcatc 480
acaaaaatcg acgctcaagt caagagggtg cgaaaaccga caggactata aagataccag 540
gcgtttcccc ctggaagctc cctcgtgcgc tctcctgttc cgac 584

```

```

<210> 640
<211> 404
<212> DNA
<213> Homo sapien

```

```

<400> 640
ccataggaac gcactcaggc aggtgggttt ttctggatgc agaaaccaga gatctagttt 60
ctatccacac agacgggaat gaacagctct ctgtgatgcg ctactcaata gatggtaoct 120
tcctggctgt aggatctcat gacaacttta ttacctcta tgtagtctct gaaaatggaa 180
gaaaatatag gagatatgga aggtgcactg gacattccag ctacatcaca caccttgact 240
ggtccccaga caacaagtat ataatgtcta actcgggaga ctatgaaata ttgtactggg 300

```

acattccaaa tggctgcaaa ctaatcagga atcgatcgga ttgtaaggac attgattgga 360
cgacatatac ctgtgtgcta ggatttcaag tatttggtgt ctgg 404

<210> 641
<211> 138
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(138)
<223> n = A,T,C or G

<400> 641
ctgtgacagg aacattacct gaagtgcagg gtggttacct gcacaaagtc ccatttccaa 60
aaatttctgt gtaattcacc agaaattttg gatggaataa ttagaaaaaa aaaaagaggt 120
taaaacntgt aactcaaa 138

<210> 642
<211> 381
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(381)
<223> n = A,T,C or G

<400> 642
ctgtagggtg aatttttacc cagaaaagat aggccctaga agcctcattt cttttctcca 60
tggaaaagga cagccctctg ctgcagcgtt caacttggtg gtttactgac agagtgaact 120
acagaaatag cttttcttcc taaaggggat tgttctacat tttgaagtta ttttttaata 180
aaattgaatt atgttggtga ttgtgcttcc taataggaaa tgcattattg gactgttttt 240
gtaacatcct gtttattgca aatagctagt atcgttcaaa aactgtataa aatacttttg 300
tacatattag caatgtctaa tttgtataca cttcagttaa atttccttaa aacttgaaag 360
gggaccttgt anaaattaaa a 381

<210> 643
<211> 403
<212> DNA
<213> Homo sapien

<400> 643
ccttcctaaa aaatagtggg gagctggagg ctacttccgc cttcttagcg tctgggcaga 60
gagctgatgg atatccatt tggccccgac aagatgacat agatttgcaa aaagatgatg 120
aggataccag agaggcattg gtcaaaaaat ttggtgctca gaatgtagct cggaggattg 180
aatttcgaaa gaaataattg gcaagataat gagaaaaagaa aaaagtcatt gtaggtgagg 240
tggttaaaaa aaattgtgac caatgaactt tagagagttc ttgcattgga actggcactt 300
attttctgac catcgctgct gttgctctgt gagtcctaga tttttgtagc caagcagagt 360
tgtagagggg gataaaaaa aaagaaattg gatgtattta cag 403

<210> 644
<211> 688
<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(688)

<223> n = A,T,C or G

<400> 644

| | | | | | | |
|------------|------------|-------------|-------------|-------------|-------------|-----|
| cctatttatt | tgttttggcc | ctggatcttt | cctaatacaca | attatatattc | tttatttttg | 60 |
| cctttgagca | gtttcatтта | tctttgtggg | caggggaagat | taaatatgaa | attcagtgcca | 120 |
| gtcattttgc | tactggttag | cttttagtttg | aggcaagtaa | aaatttttga | ttaaaattag | 180 |
| tttcttaaaa | ttatgccctt | gctttaccaa | ataatcaaат | tggctaаааа | ataagggtat | 240 |
| gtaactttgc | atттtgaaga | acaaaccaat | aatttttcat | gagccctact | cgatcttctt | 300 |
| taaagaagac | cttcctaaga | gacaattagg | gatgagtttg | attaatggga | aatagctcta | 360 |
| ggttagatta | ttttaaatтc | catacaccaa | gtgattttaac | cacagtggca | gtggcagctt | 420 |
| ctgaaccgtc | aagtatgaac | atcacttaaa | aattaaaaga | tgcttaataa | taaactctta | 480 |
| atттtcatта | agccaatctg | taattcagaa | gaaaagcata | tgtctgccat | gggactattg | 540 |
| cagtgcgtct | ccatcagtgt | taacacagga | gagatatgtt | atттtatgtg | tatgtcttag | 600 |
| tttgggatat | gtggtagtaa | gaacatgtca | agagtgcттt | tcttcaaacc | tgncagctca | 660 |
| actgangaaa | gacaggtact | tccattgc | | | | 688 |

<210> 645

<211> 484

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(484)

<223> n = A,T,C or G

<400> 645

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaaatgtgt | ctccagccca | cacttccagg | tggcagagcg | agctctctat | tactggaata | 60 |
| atgaatacat | catgagttta | atcagtгaca | acgcagcgaa | gattctgccc | atcatgtttc | 120 |
| cttccttgta | ccgcaactca | aagacccatt | ggaacaagac | aatacatggc | ttgatataca | 180 |
| acgccctgaa | gctcttcatg | gagatgaacc | aaaagctatt | tgatgactgt | acacaacagt | 240 |
| tcaaagcaga | gaaactaaaa | gagaagctaa | aaatgaaaga | acgggaagaa | gcatgggtta | 300 |
| aaatagaaaa | tctagccaaa | gccaatcccc | aggtactaaa | aaagagaata | acatgaaaac | 360 |
| gcccaggggt | acttgaatgt | ttttataaga | taggaatata | tgtcttcacc | atgggggggg | 420 |
| gtctcggatt | tcactaacgt | tgtatatgaa | aatgggtgcn | ataaaaagta | cttttaaact | 480 |
| ttgt | | | | | | 484 |

<210> 646

<211> 447

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(447)

<223> n = A,T,C or G

<400> 646

| | | | | | | |
|------------|------------|------------|------------|------------|------------|----|
| gggtcgcgtt | gaacaacttg | gttcaagatg | gtgggggcat | ttttagagcg | gcaataattg | 60 |
|------------|------------|------------|------------|------------|------------|----|


```

aaaaaaaaagg cgaactctgc cttggagagg tagatgataa gaaataaaaa ggtgtttata 120
actattttgt attataaagt gggccttaga gataggaaga agaatzgatgg attccttttg 180
gatcaatcag aaaggaaaca cgaaagaaaa gtcaggaagg tagagagaga aaaagggagg 240
gaaggagaaa gaatgggaat aaaataagga ggtaagagat actatttttg ctgagcaacc 300
agtgtgtttc aggatgatac aaagaaaaat atagaataga aataagtgca ggcttggaat 360
cagctacaaa tcctaaagat ggggtgtgtg tggatgtgtg tgtgtgtgtg tgnacaccat 420
tgtgtgtttg taaaatgtgt atgtccc 447

```

```

<210> 647
<211> 388
<212> DNA
<213> Homo sapien

```

```

<400> 647
gaagggtgata taaaatgact gtcatcattt ggagtgtgca gtacagttac ttcattgttcc 60
tcagggtttag aacaattttcc cctgcaagtt ctcacacaga taggcagaaa tcataactaa 120
ttttggttaa tcactatggc agccgttgaa gaatttaaga gaacctgcc aataagattg 180
gaataagatt ctatattatt gcatccacag aaaagaatgt actgatatac tataaactct 240
aggagaaaaac ttaattgaaa tagtgttatt aagtgttgaa agtaccataa aaatataagg 300
gaaaataagc tttcctagaa tttttcagtg ttctagtttt taaacagtga tgttttttat 360
taacctattt catccattca aagacagg 388

```

```

<210> 648
<211> 632
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(632)
<223> n = A,T,C or G

```

```

<400> 648
cctggctggg cntttgacct gcgnttttaa atnactcaca gaggggtggga caggaggaag 60
agtgaaggaa aaggtcaaac ctgttttaag ggcaacctgc ctttgttctg aattggtctt 120
aagaacatta ccagctccag gtttaaattg ttcagtttca tgcagttcca atagctgac 180
attgttgaga tgaggacaaa atcctttgtc ctactagtt tgctttacat ttttgaaaag 240
tattattttt gtccaagtgc ttatcaacta aaccttgtgt taggtaagaa tggaaattat 300
taagtgaatc agtgtgacct ttcttgatc aagattatct taaagctgaa gccaaaatat 360
gcttcaaaag aagaggactt tattgttcat ttagttcat acattcaaag catctgaact 420
gtagtttcta tagcaagcca attacatcca taagtggaga aggaaataga tagatgtcaa 480
agnatgattg gtggagggag caagggtgaa gataatctgg ggttgaaatt ttctagttnt 540
cattccgtac attttttagtt agacatcaga tttgaaatat taatgttacc tcctcaatgg 600
ggtggtatca gacctgcccg ggcgncgcn tc 632

```

```

<210> 649
<211> 300
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(300)
<223> n = A,T,C or G

```

<400> 649
 nggtgaagat agaanaaata taagcgaaat tggataaaat agcactgaaa aaatgaggaa 60
 attattggta accaatttat tttaaaagcc catcaattta atttctggtg gtgcagaagt 120
 tagaaggtaa agcttgagaa gatgaggggtg tttacgtaga ccagaaccaa tttagaagaa 180
 tacttgaagc tagaagggga agttgggttaa aaatcacatc aaaaagctac taaaaggact 240
 ggtgtaattt aaaaaaaact aaggcagaag gctttggaag agttagaaga atttggaagg 300

<210> 650
 <211> 498
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(498)
 <223> n = A,T,C or G

<400> 650
 ngtnctgnta aacagaaggg tacaangccc ttctggcttt aagcagtcac aggaatgtga 60
 cagacattcc tcttagggag cgcctcctcc taggggttcc tcatctgtct cacactgagt 120
 ggatgtaatg ctattttaat cctgctgtgg cccccaatac tagtacttgt ccataccttc 180
 ttgcattttt agcgtctgct ctgtgggggtt gttaggccct ggcactccca ggaactagtg 240
 ctaaagctgc atctntctct cccctctagg gatcgataaa gtttccactgc agaaagtctc 300
 cactgcggta tgctgacatc tgcctgaac cttcacccta cagcattaca ggctttaatc 360
 agattctgct ggaaagacac aggetgatcc acgtgacctc ttctgccttc actgggctgg 420
 ggtgatcctt ggtgcctttg tttccacaag gccttttccct gccccctgcc ttgccaaaga 480
 catttaataca gcacacag 498

<210> 651
 <211> 654
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(654)
 <223> n = A,T,C or G

<400> 651
 ctgaggggtcc ccagggtttct aaagctctca ggacgagaaa gtaggtccca agataaggag 60
 cctaaagggc ttttttcttt ctgtgtattc cttcttggcc tccaacatgg gtacagtcac 120
 aagagcatgt aacagagaag aaggactana cctaccattt tctggataaa gaattggaaa 180
 gaggatccac aggtaaccaa aaagtaccag ggaaatggca gagaaggaaa acctcaggag 240
 accaacctca taagtggat ttattagncc ctgggctcaa atccaaattg tacatgaata 300
 tgtctgggtcc tagatagggt accgaagact ttgaaagtga attttggtat atcattgccc 360
 agattccaga ctggnatttg tgtgacacaa catacaggat atatctgaat agtgctcaga 420
 agagtttgaa atgtcaaatg atattaaaat aaagattgaaa aagagaaaagc tggtcagaac 480
 ttgtggacat aacccttctg gatctgtngc ctgattaaaa aatagttgat attctcgaat 540
 gaattaaaac aagatttaga gactgagcat ggtagctnat tcttgtaatc caacnctttg 600
 ggagggcaag gcaanagaat tgcttgccgc caggagtttt gagaccagct tggg 654

<210> 652
 <211> 293

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(293)
<223> n = A,T,C or G

<400> 652
ngtctgttgc actgaggtga ctaaggatac attttgagga agtagctcca agaacatttc 60
cattttcact gtgccttcac atacatctaa tggaaatgaa cagcaccctt catccatcca 120
cggaagcgat taagaaaagg gtgggatgga aaaattaacc caacaatatt agatcaatac 180
gtagtattta agngtccata atgtgccagg ctgaagatgc acgggaaaac cacactagcc 240
ggtctgtcaa gggcttgaga ataccataaa caagaaaaca gacgaaccaa ttt 293

<210> 653
<211> 294
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(294)
<223> n = A,T,C or G

<400> 653
ngtccaccac tgcagcccta catacagttg aaaaaaaatt ccattctgtt aacatttggt 60
ttataagttt tcacgcaata cacaaaaaac ccctctgcac ttcttgtaaa gaacaaaaaa 120
gatacacaac agttaagcgt aaagatcaca ggcaatagca ttcaaactg gatgtgggta 180
gagaaaggag tacctggcat gagtacctgc ttagtttgac tgaatccttg atttttaatt 240
tggcttttca tgggcccgtc acaacaccaa cgctgtgtga ggtatggtag tcag 294

<210> 654
<211> 250
<212> DNA
<213> Homo sapien

<400> 654
ctgtccttga acaagtatca atgtgtttat gaaaggaaga tctaaatcag acaggagttg 60
gtctacatag tagtaatcca ttgttggaat ggaacccttg ctatagtagt gacaaagtga 120
aaggaaattt aggaggcata ggccatttca ggcagcataa gtaatctcct gtcctttggc 180
agaagctcct ttagattggg atagattcca aataaagaat ctagaaatag gagaagattt 240
aattatgagg 250

<210> 655
<211> 494
<212> DNA
<213> Homo sapien

<400> 655
ccattataat tttataaac cattaccctt taaattctac cgattataag cagcgtaaaa 60
gtaactatat aaagcaaaca tcgcaaagga actctgcagg agctcttaat tcctttatgt 120
agctatcata aaattcactt tcctgaagac atttactctc attcacttcc aaactccaaa 180
cctttttctg gtagcaccac ttttgttttt aatagaaaga tgagttcata tctgtacatc 240


```

cctgaaaaga aagntgctct tatggactct tgcattgttaa gactatgtct tcacatcatg      60
gtgcaaatca catgtaccca atgactccgg ctttgacaca acaccttacc atcatcatgc      120
catgatggct tccacaaagc attaaacctg gtaaccagag attactgggtg gctccagcgt      180
tgtagatgt tcatgaaatg tgaccacctc tcaatcacct ttgagggcta aagagtagca      240
catcaaaagg actccaaaat cccataccca actcttaaga gatttgcctt ggtacttcag      300
aaagaatttt catgagtgtt cttaattggc tggaaaagca ccag                          344

```

```

<210> 659
<211> 230
<212> DNA
<213> Homo sapien

```

```

<400> 659
ctgctttccc tgctaaacag ttccagagca aaagcagcaa aaagaaaata tgggagggat      60
atgggcaacg tatactcgaa cgtacgcaga gaagagagta cggtttagctc taatatttct      120
cattgaactt ggtggtatgt gccttccctg catataaggc catagtgtctt ttttgggagc      180
gctagaatat ccatccactt gacagtgtacc acaaaatagg ctgtttccag                230

```

```

<210> 660
<211> 80
<212> DNA
<213> Homo sapien

```

```

<400> 660
ctggtccttg ttaaaactga tcaccacttt ggagagatcg actggagggt cctgggtgtt      60
ctgagggggc tgggggacag                          80

```

```

<210> 661
<211> 535
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (535)
<223> n = A,T,C or G

```

```

<400> 661
ctgaaccata tctgattaac tctttggtct ctgttatttg aacaaaaccg acgctatgcc      60
tgcagccgcc agactgcaac caaaaacaca gtttggggtc agaagacatt aaaaatcaca      120
ataaaatagg atgaatgttc taagtcacgc aactgaatca aggcaccttt ttttttcaaa      180
agcaaaaagt tgtttaacaa tattccagaa tagtagatac ttcaaaaacc agattacagt      240
atatatcatt ttgctgcaca ttttagtcta ttttctgtat acatagtcac acattcttta      300
ccctctccca acttatacat gctttatccc cccagtcatg tgctatgtag gtataaaaaa      360
ataaagtgtg atctaaacaa gtgattttaa aaaaaaaact aacgaatgcc ncnatnataa      420
cnctgaactt gtttcctnt tgaaggacat tggaaatgtt accgaggttn ntttacctng      480
gccgcaaccn cnctangggc naattccagc ncactggggg ccgttactag gggat          535

```

```

<210> 662
<211> 257
<212> DNA
<213> Homo sapien

```

```

<400> 662

```

```

cctgactaaa gcacatatca cactccctac acttccatgt tttctctccc atgtggaccc      60
tctgatgcat atcaagattc aagcgctgtg tgtagccctt cccacagtcc tcacatttgt      120
atggcctttc tacactgtga actttttctt gcactttaga gaatgaattc tgtacaatgt      180
tcttcccatg ctgctcacat ttgagagggtg tttctctgct gtggcgtctc tgatgggtca      240
gacgagttga ggaccag                                     257

```

```

<210> 663
<211> 516
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(516)
<223> n = A,T,C or G

```

```

<400> 663
ccaattatag gtatttttatt ttttaaagat tagagngttc ttgaagctct ttctatttct      60
ttgtcaatga actaaacatt ggcaaatatg tagggtttcc cacataagaa cattattaac      120
atcaaaatag aaagctggtg gtagaaataa tgattgggaa cacagagtct ctactcagcg      180
ttctacttct gccataccat aactttgtga tctcacgaaa tatctctcca tgttctcctc      240
cctatgtata gttctgtcat ttttcaataa gagctttttg ctttaattatg aagtactagt      300
tactataacc attattttga gcttcatgta aatcaagaac acatggactc cacttgcaaa      360
acattgaaaa tgtagttagg gattgggggc aaaaagcaac atttttaaatt gtgtaaagac      420
aatgagtaag caacaaagtg tccaattttt taggcgaaaag ttgcatatgt caggaaaagg      480
caggattaag taatagagaa tttgaatgat aactgg                                     516

```

```

<210> 664
<211> 212
<212> DNA
<213> Homo sapien

```

```

<400> 664
gtccgaggag gttagtgtgt gcaataaaaa tgattaagga tactagtata agagatcagg      60
ttcgctcttt agtggtgtgt atggctatca tttgttttga gggttagtttg attagtcatt      120
gttgggtggt aattagtcgg ttgttgatga gatatttgga ggtggggatc aatagagggg      180
gaaatagaat gatcagtact gcggcgggta gg                                     212

```

```

<210> 665
<211> 408
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(408)
<223> n = A,T,C or G

```

```

<400> 665
atccaggggt ncccggtngc tgcngggaaa cctccagcct tgttcttcaa accactcagc      60
tcatgtgttt tgcgctgact agtactgaat aatacaacca ctcttattta atgttagtat      120
tatttatattg acaactcagt gtctaacagc ttgatatgca ggtccttgca tcttacattt      180
cttttaggaag ttaccatttt gtaactttta aaacaggaaa aatatcagtt ggcaaatgca      240
atcttttttt tttttaagct aaaggggggn naacngnaan naaaatnttt ntgangtnng      300

```

gtctataagc acccttggang ggatntgtta aaagnngncat naanggggga ttctcntttt 360
gcaaaaaaat ntaannatca atttatanan ctttatTTTT nactttnt 408

<210> 666
<211> 635
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(635)
<223> n = A,T,C or G

<400> 666
ctgaagnaca agggtcaggc aaaaataaga tcacaatcac caatgaccag aatcgcttga 60
cacctgaaga aatcgaaagg atgggttaatg atgctgagaa gtttgctgag gaagacaaaa 120
agctcaagga gcgcattgat actagaaatg agttggaaag ctatgcctat tctctaaaga 180
atcagattgg agataaagaa aagctgggag gttaaaccctc ctctgaagat aaggagacca 240
tggaaaaaag tgtagaagaa aagattgaat ggctggaaag ccaccaagat gctgacattg 300
aagacttcaa agctaagaag aaggaaactgg aagaaattgt tcaaccaatt atcagcaaac 360
tctatggaag tgcaggccct cccccaactg gtgaagagga tacagcagaa aaagatgagt 420
tgtagacact gatctgctag tgctgtaata ttgtaaatac tggactcagg aacttttggt 480
aggaaaaaat tgaaagaact tanctctcga atgtcattgg aatcttcacc tcacagtggg 540
gttgaaactg ctatagccta agcnggctgt ttactgnttt ncattagcag gtgctcacca 600
tgtctttggg gtgggnnggg ggagaaagaa agaan 635

<210> 667
<211> 388
<212> DNA
<213> Homo sapien

<400> 667
gaagggtgata taaaatgact gtcattcattt ggagtgtgca gtacagttac ttcattgttcc 60
tcagggtttag aacaattttcc cctgtaagtt ctacacacaga taggcagaaa tcataactaa 120
ttttgggttaa tcaactatggc agccgttgaa gaatttaaga gaacctgcca gtaagatttg 180
gaataagatt ctatattatt gcatccacag aaaagaatgt actgatatac tataaactct 240
aggagaaaaac ttaattgaaa tagtgttatt aagtgttgaa agtaccataa aaatataagg 300
gaaaataagc tttcctagaa tttttcagtg ttctagtgtt taaacagtga tgttttttat 360
taacctattt catccattca aagacagg 388

<210> 668
<211> 498
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(498)
<223> n = A,T,C or G

<400> 668
tgatcttaac aaaattcgta gcagtggaaac cttgaaatgc atgtggctag atttatgcta 60
aaatgattct cagtttagcat tttagtaaca cttcaaagg ttttttttgt ttgttttcta 120
gacttaataa aagcttagga ttaattagaa gaagcaatct agttaaattt cccatttgta 180

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ttttattttc | ttgaataactt | ttttcatagt | tattcgttta | aaaagattta | aaaatcattg | 240 |
| cactttggtc | agaaaaataa | taaatatatc | ttatgaatgt | ttgattccct | tccttgctat | 300 |
| ttttattcag | tagatttttg | tttggcatca | tgttgaagca | ccgaaagata | aatgatTTTT | 360 |
| aaaaggctat | agagtccaaa | ggaatgttct | tttacaccaa | ttcttccttt | aaaaatntct | 420 |
| gaggaatttg | ttttcgctt | actttttttt | cttctgtcac | aatgctaagn | ggtatccgag | 480 |
| gtntttaata | tgagattt | | | | | 498 |

<210> 669
 <211> 622
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| <400> 669 | | | | | | |
| ccttagccaa | agaatgcagt | ggagccttcc | cccttcaact | gcattgtgaa | tgaataccaa | 60 |
| ttaacagcat | aaaaattaat | agtcccatat | cagatctgga | aggggtttct | ggggctgtct | 120 |
| gatgtcccta | tcctgttgta | gtgaacacaa | tagcagaaaa | ttctttctgg | gtccatctgc | 180 |
| tataaagtct | tggtaaaaca | gcattactat | gaagaggatg | aactcaccta | ccttcagatg | 240 |
| gaggaaaagt | gaaaaggact | taggcttttag | tcctccatga | cttttcttaa | gcactaccta | 300 |
| cctgtaataa | gctgagtgc | aaaggatgcc | gaagaaaatc | tgacccaga | agctgttaga | 360 |
| aagcactgca | gagaacaggg | tatgaagaaa | ataaagagtt | cttaataaac | ccttaagatt | 420 |
| ctttgttcaa | ggtaaccttg | ccaaaagggc | agagtaggtg | gcaaagagtt | gcttttaatc | 480 |
| tagctctaca | ctgcatttga | aaataaaaatt | tgcccatttt | gaatatattg | tttataatta | 540 |
| aatgtgcttt | ttacactgca | ggtcaatata | aaaactgggt | agtaaatttc | cagcgagcat | 600 |
| ttatgttcat | ttgctcacag | ca | | | | 622 |

<210> 670
 <211> 477
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| <400> 670 | | | | | | |
| ttgggccctc | tagatgcagt | ctcgagcggc | cgccagtgtg | atggatatct | gcagaattcg | 60 |
| cccttgccgc | ccgggcaggt | gatggatgag | gagcaaaaac | tttatacggg | tgatgaagat | 120 |
| gatatctaca | aggctaataa | cattgcctat | gaagatgtgg | tcggggggaga | agactggaac | 180 |
| ccagtagagg | agaaaataga | gagtcaaacc | caggaagagg | tgagagacag | caaagagaat | 240 |
| atagaaaaaa | atgaacaaat | caacgatgag | atgaaacgct | cagggcagct | tggcatccag | 300 |
| gaagaagatc | ttcggaagaa | gagtaaagac | caactctcag | atgatgtctc | caaagtaatt | 360 |
| gcctatttga | aaaggttagt | aaatgctgca | ggaagtggga | ggttacagaa | tgggcaaaat | 420 |
| ggggaaaggg | ccaccaggct | ttttgagaaa | cctcttgatt | ctcagtctat | ttatcag | 477 |

<210> 671
 <211> 127
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| <400> 671 | | | | | | |
| gtgtgtgtgt | ctacttgggc | gtgtttaacg | tgtgcgtttg | tgtctgcgtg | tgcatgtgtc | 60 |
| tgtgtgtgcg | cgtgtatttc | agtttggtt | gccggatccc | atatgattgc | gtgcctgtgt | 120 |
| acctgag | | | | | | 127 |

<210> 672
 <211> 400
 <212> DNA
 <213> Homo sapien

<400> 672
 gggctctgcac agctatgtta acagcatcct tataccagga gtaggaggaa agacacgact 60
 ggaaaagcaa ttcaagctgg tcacacagtg taatgcaaaa tatgtggaat gtttcagtgc 120
 tcagaaagag tgtaacaaag aaaagaacag aaactcttca gttgtgccat ctgagcgtgc 180
 tcgagtgggt cttgcaccat tgcttggaaat gaaaggaaca gattacatta atgcttctta 240
 tatcatgggc tattatagga gcaatgaatt tattataact cagcatcctc tgccacatac 300
 tacgaaagat ttctggcgaa tgatttggga tcataacgca cagatcattg tcatgctgcc 360
 agacaaccag agcttggcag aagatgagtt tgtgtactgg 400

<210> 673
 <211> 600
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(600)
 <223> n = A,T,C or G

<400> 673
 ctggcggttg tcattagtga atgtatgaca gcaggatgtg aggggatgcc caggagtcag 60
 tgttagcatt gtcactgtg atcactgcta ttaatatcat ccattaattt attagtgagc 120
 ttcactatat gcagactggg agataaggag aaaatctgtc acattctctc tagctaatac 180
 gatcagctac caattaatga gattctgaat gaaatatcaa tatgtgtttt tctaatttgg 240
 acctaggaca gagctgttg tttgtcataga gaaaaacaat aatgcttaaa catagcacat 300
 tataattaaa gcagggttct cacatacttt tcattttatc ctttggataa ttttgtgagg 360
 aacgcaggac accaacttcc ctttcataga tacaatcccc atgctattga tgaaagtgtt 420
 tttgaatgaa gccatacaac aaataactga tcaaagtggc attacacca aatttcttag 480
 taggactcct gcatagaatg tttagataga cgtgaaaagt ttgttcanga ggaccagcaa 540
 gagagaaact gggttctttg ggagggtttc ggtgctacat ttataccctn catcagagtn 600

<210> 674
 <211> 140
 <212> DNA
 <213> Homo sapien

<400> 674
 ggtgggttgg gttaaatgagt gaggcaggag tccgaggagg ttagttgttg caataaaaaat 60
 gattaaggat actagtataa gagatcaggt tcgtccttta gtgttggtga tggctatcat 120
 ttgttttgag gttagtgtga 140

<210> 675
 <211> 245
 <212> DNA
 <213> Homo sapien

<400> 675
 gttgggttgg tgggtgtaaat gagtgaggca ggagtcaggag gaggttagtt gtggcaataa 60
 aaatgattaa ggatactagt ataagagatc aggttcgtcc ttagtggttg tgtatggcta 120
 tcatttgttt tgaggttagt ttgattagtc attgttgggt ggtaattagt cggttgttga 180
 tgagatattt ggaggtgggg atcaatagag ggggaaatag aatgatcagt actgcggcgg 240
 gtagg 245

<210> 676
 <211> 621
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(621)
 <223> n = A,T,C or G

<400> 676
 ctgtccccag ggnaaatagt ngaattcaac taagatctgt taataagatg tcagaataac 60
 taataatttt attaggaaaa aatcatgttt taaatttcaa aatgacactt atttgtcaag 120
 taatatgatac ttggaaaatt ttaaagaaaa ataatacctac ttataaacta cttttttata 180
 attgttttca gaaaaaaagt ttacagtctt aaggaaaata ttcagggtcta tcatatgggt 240
 tgacagattt tttaaaagt atttttggtt aggtcttctt ttagaaaaaa attaacttca 300
 aggggttttt gtaccactat aatctctaata acttactcag aattactgtg tatttactta 360
 atttcttatt atgtgcctta ttatgtgctt aagatacaat aggttagagt ttaatctaaa 420
 tatcttgaaa gctatattgt gggcttggtt agcattttgt tttttctttc tctgttttgg 480
 taaggattta aaattttttt cattgcaatt ttaagtgggt ttcaataagt aatagttttt 540
 atcaaatttt tgggtgcttgg tgcagagacg gcgtggggaa ggggtgaatgg ttttgggaat 600
 aattcagtgc acacctgggg g 621

<210> 677
 <211> 210
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(210)
 <223> n = A,T,C or G

<400> 677
 tttacataaan atattatcag catttaccat ctcacttcta ggaataactag tatatcgctc 60
 acacctcata tcttccctac tatgcctaga aggaataata ctatcactgt tcattatagc 120
 tactctcata accctcaaca cccactccct cttagccaat attgtgccta ttgccatact 180
 agtcttttggc gcttgccaag cagcggttagg 210

<210> 678
 <211> 383
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(383)
 <223> n = A,T,C or G

<400> 678
 gtaggagtca ggttagttagg gttaacgagg gtggtaagga tggggggaat tagggaagtc 60
 aggggttaggg tgggttatagt agtgtncatg gttatttagga aaatgagtag atatttgann 120
 aactgattaa tgtttggggnn tgagttnta tatcacagcc anaattntat gatgnaccat 180
 gtanogaaca atgctacagg gatgaatatt atggagaagt antctanttt gaagcttagg 240

gagagctggg ttgtttgggt tgnnggetcan tgtcagttcc anataataac ttcttgggtct 300
 aggcacatga atattgttgt ggggaanaga ctgataataa aggtggatgc gacaatggat 360
 ttacataat gggggtatna gtt 383

<210> 679
 <211> 371
 <212> DNA
 <213> Homo sapien

<400> 679
 aaaatgaaaa tattgacaag agtttcagat agaaaaatgaa aaacaagcta agacaagtat 60
 tggagaagta tagaagatag aaaaatataa agccaaaaat tggataaaat agcactgaaa 120
 aaatgaggaa attattggta accaatttat tttaaaagcc catcaattta atttctgggtg 180
 gtgcagaagt tagaaggtaa agcttgagaa gatgaggggtg tttacgtaga ccagaaccaa 240
 tttagaagaa tacttgaagc tagaagggga agttgggttaa aaatcacatc aaaaagctac 300
 taaaaggact ggtgtaatTT aaaaaaaaact aaggcagaag gcttttggaa gagttagaag 360
 aatttgggaag g 371

<210> 680
 <211> 176
 <212> DNA
 <213> Homo sapien

<400> 680
 cctaggattg tgggggcaat gaatgaagcg aacagatTTT cgttcatttt ggTtctcagg 60
 gtttggttata atTTTTtatt tttatgggct ttggtgaggg aggtaagtgg tagtttTgtgT 120
 ttaatatTTT tagttgggtg atgaggaata gtgtaaggag tatgggggta attatg 176

<210> 681
 <211> 152
 <212> DNA
 <213> Homo sapien

<400> 681
 ctggagatgg atatgagact agtcaagatg tgaatgctaa ttggagagaa atataatttt 60
 aggaagatgc acattgatgt ggggttttga tgtgtctgat tttgactact caagctctgt 120
 ttacagaaga aaattgaatg gcgaggggtg gg 152

<210> 682
 <211> 141
 <212> DNA
 <213> Homo sapien

<400> 682
 ccagtgcTtg cttgccgtgg tttagtgatt gggTgttaga aataaaaaact caggtctatt 60
 tcttaccagt cagtaacaat ttttagagaa tgtacttggt atataatata tggacttcag 120
 gaactttgtt ggggtggggg g 141

<210> 683
 <211> 308
 <212> DNA
 <213> Homo sapien

<400> 683

<210> 687
 <211> 315
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(315)
 <223> n = A,T,C or G

```

<400> 687
nngtctgtga aaaactcttt ggatgattct gccaaaaagg tactttctgga aaaatacaaaa      60
tatgtggaga attttgggtct aattgatggg cgctcacca tctgtacaat ctctgttttc      120
tttgccatag tggctttgat ttgggattat atgcaccctt ttccagagtc caaaccggtt      180
ttggctttgn gtgtcatatc ctattttgtg atgatgggga ttctgacctt ttataacctca      240
tataaggaga agagcatctt tctcgtggcc cacaggaaa atcctacagg aatggatcct      300
gatgatattt ggag                                     315
  
```

<210> 688
 <211> 522
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(522)
 <223> n = A,T,C or G

```

<400> 688
ctgaattaga ggaggagaaa agaagccatt nnggagtact ttaattgttt agatgtgaga      60
ggctgaatgt ttgggttaag atgtagttg tcagaatcat gagaaaagg ttttaagcaag      120
gggcattttc aattctaaaa ataacaacta ctgttattta ttgagcacta tctttttggt      180
gggtactgtc taaagtactt gatttatttt ttaaaaacctt acaaaaaact tacaaggtag      240
gtactgaaag attcagtaat ttgttcaaag tcacacagca aataagcaac agactctgga      300
tttgaaccag gcaatcctag agcctgtact gttagtaatt ataactttagc acctgtcaag      360
aattcctgtt gagtgtcaag aagcaancac caagttagga tttaaagcaa acatgattga      420
agaatactgt ggtgtggttg acagtagtgc ctaagtctgt tttcagagtg aaaaatgaca      480
aattagattt taagtatggt ttggagataa tatcaggaca gt                               522
  
```

<210> 689
 <211> 158
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(158)
 <223> n = A,T,C or G

```

<400> 689
tctcaactta ntntnatacc cacaccacc caanaacagg gtttgtagg nattgtttgc      60
attaataaat taaagctcca tagggctctt tcgtcttctg gtgtcatgcc cgctcttca      120
cgggcaggtc aatttcactg gttaaaagta agagacag                                     158
  
```

<210> 690
 <211> 300
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(300)
 <223> n = A,T,C or G

<400> 690
 tagaactcgt atttttaaac ttctattctc tanccttttc cactacatta tgacacaaga 60
 ccttgacagaa agtcgtctgg aaaatatcag accatctctt acttgcccc tccaatctta 120
 catcgaatta tatgcaccct taaaaagtta tttggagttt taaaaaactc tattagccca 180
 aattacctga aataaactcc tggcttggtc ccctaagtgt tataaaaaat tgattgaaaa 240
 tattcatttt aaaaatgaag ntcttgaatt tatttaaatt actgtcttgc agtgagttgg 300

<210> 691
 <211> 305
 <212> DNA
 <213> Homo sapien

<400> 691
 ctgttcagaa agctcattgg acctggtttt gaaaataaaa caaagttaaa accctggggag 60
 gagttattgt gcagtgtgga gtactcaggc tttcttataa agaaaaaaaa agttatctgg 120
 taccaaagtg tgcaacctac agaccctcag gtactgccct gtgacttctc tgtatgacat 180
 cacaaggctg ccaagtgcct gtttttctag aactaggagt tggtagaggt ttggtagtgc 240
 tgaaaccatg cataggattg gtttactaaa ttaaaacctt attacgtacg tcctccaaaa 300
 gacag 305

<210> 692
 <211> 582
 <212> DNA
 <213> Homo sapien

<400> 692
 caggaaatgg ataaccattt taactgtatt ttttgcagcc cgtaccttct tgggaataca 60
 attgtctaac tttttatttt tggctctggc gttgtggtgt gcaaaactcc gtacattgct 120
 attttgccac actgcaacac cttacagatg tggaagatgt gaaatttgtc atcaattatg 180
 actaccctaa ctctcagag gattatattc atcgaattgg aagaactgct cgcagtacca 240
 aaacaggcac agcatacact ttctttacac ctaataacat aaagcagggtg agcgacctta 300
 tctctgtgct tcgtgaagct aatcaagcaa ttaatcccaa gttgcttcag ttggtcgaag 360
 acagaggtgc aggttaaggat gactgatagg aaatgttggg agttacgagt cacatcggtg 420
 tctacaaatc catttaaatg gtattggagg gtgagtaaaa ccttgaatgt gaaaaacttaa 480
 gctgaaaaat tgtaaaaaaca tttcacgcct accatgaata gatctgtttc tttctgtcca 540
 caatgatttg tgtcatagac ataattgatc aatttgaat tg 582

<210> 693
 <211> 275
 <212> DNA
 <213> Homo sapien

<400> 693

```

ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt      60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct      120
atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg      180
gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaagggtg      240
ataagctctt ctatgatagg ggaagtagcg tcttg                                     275

```

```

<210> 694
<211> 397
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(397)
<223> n = A,T,C or G

```

```

<400> 694
nggtctgcat ttttattgcg atctgcagat gaactggaaa atctcatttt acaacagaac      60
tgagacagac gaccaccata ttactgagg tctaaatttg cagtttccac taatgacatt      120
ttgatttccc aacagagata cttctggtct tactgcacag tcttttaaga gaaatacttc      180
cattatgcca cattgtcctt gatccgtaag tgatgtgta aggtgcttca aaggaactct      240
gacctctgaa gtacttgagc tacttttagta tgtccagcct attgcttttt gtttttagtgt      300
gtcaccataa atatcagggg cataaaaaggc tatctattct taattcaagg ataaaaacaga      360
agaagcttgt ggtataaaac aatagttcaa gatccag                                     397

```

```

<210> 695
<211> 609
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(609)
<223> n = A,T,C or G

```

```

<400> 695
ctgagcttcc atttgtcagc tagcactgng gtagtcaacc atgcgaatga ggctattttg      60
gacctcatga ttgtccagtg cctgggctga taccngggga aacgaaattt tgtggctgcc      120
cacaaaatca tggaaaataa tgatttttta gaaaacctcc actgntttgt tgtgcagcaa      180
taaataactg aaacaccaat ccaaaaaact tataaagcta taacaattaa aacagnataa      240
taatagtncc gggatacaaa aatgggtcaaa ttgaagagga tacaaaagcct caaagcagtc      300
ctcactcata ananccttgt tgtatcacta aaanggcatt aaaattgaga anaaggaana      360
actagtggat taattaataa atgagaagta tccataagga aaaattaataa ttnnattctt      420
gcttcacatt atgaaaaaat acaaacaaca gattgattaa agacttaaat gngatcaaca      480
aaatgttaaa actgtgataa gaacatttaa gaaaatagtt ctatnaccct gggataaaac      540
attttcntcc aaggcattaa agtgttaaat gaaaagactg atncatttat tcattagaat      600
ttaaattcn                                     609

```

```

<210> 696
<211> 300
<212> DNA
<213> Homo sapien

```

```

<400> 696

```

```
<210> 697
<211> 391
<212> DNA
<213> Homo sapien
```

<400> 697

```
<210> 698
<211> 536
<212> DNA
<213> Homo sapien
```

<400> 698

```
<210> 699
<211> 419
<212> DNA
<213> Homo sapien
```

```
<220>  
<221> misc_feature  
<222> (1) ... (419)
```


<223> n = A,T,C or G

<400> 699

| | | | | | | |
|-------------|------------|------------|-------------|-------------|------------|-----|
| ngtccacctg | agggcaggtg | acaaggacct | gacagagccc | atgcagggct | ttagatttgg | 60 |
| acacacaaga | gttgataact | tcctcatgaa | ctccttgcct | gatctaaact | catattatgg | 120 |
| gttctgactg | tttgagtaat | catcttcaag | gttaaaccctc | ttggcagtta | cccttttcac | 180 |
| aaagtgcaca | gtgggaatcg | agaatcgata | gggttaattt | tggagcagtg | gcttatacca | 240 |
| ttcacctctg | tttttttgtg | attatttcac | agataatgag | accttaataa | caaataggcg | 300 |
| taaaaaaatt | ttcacattga | aatgatagaa | acatttgatg | taataaaaact | tggttggtt | 360 |
| gatatttttaa | ggaattgaaa | cctagcaatc | ttattggaga | gacaagaatt | ggtctccag | 419 |

<210> 700

<211> 336

<212> DNA

<213> Homo sapien

<400> 700

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccacttattg | tccttaaaaa | tccatactga | tacatggaca | gtaagtgtgt | tttcagatgg | 60 |
| agtaccagca | ccgaaaatgg | gttgagggag | gatgggttgt | atgtatgttt | ctgcccacta | 120 |
| attttgagca | gccatattat | gaattaaatc | gtcacagcca | agtaataacc | caagaatggg | 180 |
| atgagtttca | tgtgtaatag | ctcaaattga | ataagcatga | atgctggagt | ggaccattat | 240 |
| cctcaaatat | tctatgtcac | ttctcattta | aagactcttg | ttatgaacta | ttagaaactt | 300 |
| taggcaaaat | caaaagtatt | tgcggcacaa | ttaaagg | | | 336 |

<210> 701

<211> 418

<212> DNA

<213> Homo sapien

<400> 701

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccatgtgatg | atgttgacaa | cccctgaaga | gcctcagtc | attgttccac | gtttaagaac | 60 |
| taggaatacc | aggactgatg | caattctact | gggtcactat | cgcttgtcac | aagacacaga | 120 |
| caatcagacc | aaagtatttg | ctgtaataac | taagaaaaaa | gaagaaaaac | cacttgacta | 180 |
| taaatacaga | tattttcgtc | gtgtccctgt | acaagaagca | gatcagagtt | ttcatgtggg | 240 |
| gctacagcta | tgttccagtg | gtcaccagag | gttcaacaaa | ctcatctgga | tacatcatte | 300 |
| ttgtcacatt | acttacaaat | caactggtga | gactgcagtc | agtgtttttg | agattgacaa | 360 |
| gatgtacacc | cccttgttct | tcgccagagt | aaggagctac | acagctttct | cagaaagg | 418 |

<210> 702

<211> 261

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(261)

<223> n = A,T,C or G

<400> 702

| | | | | | | |
|-------------|------------|-------------|------------|------------|-------------|-----|
| gggcctgttg | tgggggtggg | ggaagcaggg | aggggaacag | ctaaataggt | tgctgttgat | 60 |
| ttgggttaaaa | aatagttagg | ggatgatgct | aataattagg | ctgnnggtgg | ttgtgttgat | 120 |
| tcaaattatg | tgttttttgg | agagtcagtg | cagtggtaga | aatataattg | ttgggaacnat | 180 |
| tagnttttagc | attggagtag | gtttagggtta | tgtacgtagt | ctaggccata | tgtgttggan | 240 |
| attgagacta | gtagggctag | g | | | | 261 |


```
<220>  
<221> misc_feature  
<222> (1)...(266)  
<223> n = A,T,C or G
```

```
<210> 707
<211> 358
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(358)
<223> n = A,T,C or G
```

```
<210> 708
<211> 491
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(491)
<223> n = A,T,C or G
```

| | | | | | | | |
|-------------|------------|-------------|-------------|------------|------------|--|-----|
| <400> | 708 | | | | | | |
| cctactatgg | gngttaaatt | ttttactctc | tctacaaggt | tttttcctag | tgtccaaaga | | 60 |
| gctgttcctc | tttggactaa | cagttaaatt | tacaagggga | tttagagggt | tctgtgggca | | 120 |
| aattttaaagt | tgaactaaga | ttctatcttg | gacaaccagc | tatcaccagg | ctcggtaggt | | 180 |
| ttgtcgccctc | tacctataaa | tcttccact | atthtgcctac | atagacgggt | gtgctctttt | | 240 |
| agctgttctt | aggtagctcg | tctggtttctg | ggggctcttag | ctttggctct | ccttgcaaag | | 300 |
| ttattttctag | ttaattcatt | atgcagaagg | tataggggtt | agtccttget | atattatgct | | 360 |
| tggttataat | ttttcatctt | tcccttgcg | tactatatct | attgcgccag | gtttcaattt | | 420 |
| ctatcgcccta | tactttattt | gggtaaatgg | tttggctaag | gttgtctggg | agtaagggng | | 480 |
| gagtgggttt | g | | | | | | 491 |

<210> 709
 <211> 460
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(460)
 <223> n = A,T,C or G

<400> 709
 nggttttttt ttagagcaa ataatttatg caaaatatgt tacaaaatct gggatgctaa 60
 atagttgaca caagtactgt gtttgacatt tagtttcatt tgaattagta atagaatttg 120
 ctcttccaa catttacatc ttttttcttt ctgactttat atattttcaa taaaaatttg 180
 ctccacagtt ttttaagntca ttcttcttga atccgntttt acatttgctg ngacaaacct 240
 gcataaaact agattttata gatataactt ctttggaaga gataaaaatt caaaagtttg 300
 acattgcttt canttattct tttcttcatt gttttgattg gcccctgtta gattgatgta 360
 ttgccaatct acttttgatg gcatgaatnt aaaatgacaa cataaaaagc ncttctagtg 420
 caacagtaat tgaaacttgc agttttccat taaaaaaaaa 460

<210> 710
 <211> 542
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(542)
 <223> n = A,T,C or G

<400> 710
 ctgttacagt gacaagagat aaaaagatag acctgcagaa aaaacaaact caaagaaatg 60
 tgttcagatg taatgtaatt ggagtgaata actgtgggaa aagtggagtt cttcaggctc 120
 ttcttggaag aaacttaatg aggcagaaga aaattcgtga agatcataga tctactatg 180
 cgattaacac tgtttatgta tatggacaag agaaatactt gttgttgcag gatatactcag 240
 aatcggaatt tctaactgaa gctgaaatca tttgngatgt tgtatgcctg gtatataatg 300
 tcagcaatcc caaatccttt gaatactgtg ccaggatttt taagcaacac tttatggaca 360
 gcagaatacc ttgcttaatc gtagctgcaa agtcagacct gcatgaagtt aaacaagaat 420
 acagtatttc acctactgat ttctgcagga aacacaaaat gcctccacca caagccttca 480
 cttgcaatac tgctgatgcc cccagtnagg atatctttgt taaattgaca acaatggacc 540
 tg 542

<210> 711
 <211> 394
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(394)
 <223> n = A,T,C or G

<400> 711
 caaacccact ccaccttact accagacaac cttagccaaa ccatttacct aaataaagta 60

```

taggcgatag aaattgaaac ctggcgcaat agatatagta ccgcaaggga aagatgaaaa 120
attataacca agcataatat agcaaggact aacccctata ctttctgcat aatgaattaa 180
ctanaaataa ctttgcaagg agagccaaag ctaagacccc cgaaaccaga cgagctacct 240
aagaacagct aaaagagcac acccgtctat gtagcaaaat agtgggaaga tttataggna 300
gaggcgacaa acctaccgag cctggtgata gctggttgtc caagatagaa tcttagttca 360
actttaaaatt tgcccacaga accctctaaa tccc 394

```

```

<210> 712
<211> 552
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(552)
<223> n = A,T,C or G

```

```

<400> 712
gaggctctgta naatgccagg ctcaaatttg tctttataat ttaataccag aaatctttcc 60
cttgtgatgt ttctttcttt ctggattgcc tctatagcag gggatagcgg gggaggataa 120
ggcacatctt tgntgtactg agaaatttga ccacgcagga tgatgtggct gttctcattc 180
atctgcacag agaaaaataa tgataaaata tccctttcct atgtttactg attttatggc 240
tgccataatg gaagcctcct tgactattta atcctttctg tcaactaggt tcgatttttt 300
ttttaattta cctgtagtag gtatttaana attttaacta gctanaaata attacattcc 360
aaaggaacac caaggcaaat aaatggttg taatcagcaa aagaattaca ttagttgttg 420
ntgctactta ttagggggag aactgttttt ttttaaattt aaacaattta ataatctcaa 480
ctgcaaataa ttttagatgc agcaaaggac tatgtagncg ttaatacctc atgttgatat 540
tttcataata tt 552

```

```

<210> 713
<211> 518
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(518)
<223> n = A,T,C or G

```

```

<400> 713
ccaaaaactg gaagcagctc actaaacaaa cagtggcata cccatagaac tgcatacttc 60
tcagcagtat gaaagaatga gctacttata taagcatcat tgataaacct caaaaaaaaa 120
atgccacatg aanaaaccca aagggganaa acataaaaaac tttatatgtc agtcatataa 180
aattctanaa aatgcaaact aatccatcnt aaaggaaaagt aaatcaacag ttgtctggag 240
gaccananag agcaggagga ganagattat taaaggggtt aaagtaaatt tgggagtgcc 300
cttcnctttt taaatnctat gaaaatgaaa gtaaaggcnc atgcatgttg taaactaata 360
gtaacaaaca naatgggttg gagtgggttg ttgtctgggg acatcattac aaaatgtaag 420
ccagtttatn taaattttga aaagaccgtg gactctgatc tgactgatna atgttggaag 480
agataagtgt gctgcaaagt ggggaattaa taaaacag 518

```

```

<210> 714
<211> 281
<212> DNA
<213> Homo sapien

```

```
<210> 715
<211> 443
<212> DNA
<213> Homo sapien
```

```
<210> 716
<211> 639
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(639)
<223> n = A,T,C or G
```

```
<210> 717
<211> 473
<212> DNA
<213> Homo sapien
```

```
<220>  
<221> misc_feature  
<222> (1)...(473)
```

<223> n = A,T,C or G

<400> 717

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| nntgaggcta | ctgctgtttt | attacaacat | tacctcttgt | ttttataaag | tgtaccaaga | 60 |
| tttaaattga | taactttatt | ttacttgaaa | aaaaaaagtt | tnntttatca | ccagtgttac | 120 |
| agttgtcttc | tgtttctttt | tgttttgntt | tatttgnttt | cctttttagc | caaagagtga | 180 |
| acagaanatt | ttcttatttt | ggtggctatt | cattttactt | ttaaaagtga | ttggtggatt | 240 |
| ttagactaat | tatgggggaa | tttgccacca | aaataaaaaa | tatgtaaagn | gtagtgatta | 300 |
| cagagtgggt | aaaatgtggg | ttagtactta | tttattccat | taattgatta | tttgactgtt | 360 |
| tataaagaaa | gttgctttat | ttcttttaac | atcttcaaaa | gatgatcctt | tcttgtcaca | 420 |
| ttatagccaa | aagaagcaga | gaacttcact | gtctgcattt | ggttcctggt | tgg | 473 |

<210> 718

<211> 207

<212> DNA

<213> Homo sapien

<400> 718

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ggtaaagtct | agtataatat | ttaccatctc | acttctagga | atactagtat | atcgctcaca | 60 |
| cctcatatcc | tccttactat | gcctagaagg | aataatacta | tcactgttca | ttatagctac | 120 |
| tctcataacc | ctcaacaccc | actcctctt | agccaatatt | gtgcctattg | ccatactagt | 180 |
| ctttgccgcc | tgccaagcag | cggtagg | | | | 207 |

<210> 719

<211> 255

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(255)

<223> n = A,T,C or G

<400> 719

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| cctatattac | ggatcatttc | tctactcaga | aacctgaaac | atcggcatta | tcctcctgct | 60 |
| tgcaactata | gcaacagcct | tcataggcta | tgtcctcccg | tgaggccaaa | tatcattctg | 120 |
| agggggccaca | gtaattacaa | acttactatc | cgccatccca | tacattggga | cagacctagt | 180 |
| tcaatgaatc | tgaggaggct | actcagtaga | cagncccacc | ctcacacgat | tctttacctt | 240 |
| tcacttcate | ttgcc | | | | | 255 |

<210> 720

<211> 455

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(455)

<223> n = A,T,C or G

<400> 720

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaatgtcga | aacctacaag | atttccttaa | aatctctaat | agaggcatta | cttgctttca | 60 |
| attgacaaat | gatgccctct | gactagtaga | tttctatgat | ccttttttgt | cattttatga | 120 |
| atatcattga | ttttataatt | ggtgctattt | gaanaaaaaa | atgtacattt | attcatagat | 180 |

```

agataagtat caggtctgac cccagtggaa aacaaagcca aacaaaactg aaccacaaaa      240
aaaaaggctg gtgttcacca aaaccaaact tgttcattta gataatttga aaaagctcca      300
tagaaaaggc gtgcagtact aagggaacaa tccatgtgat taatgnttnc attatgttca      360
tgtaanaagc cccttatttt tagccataat tttgcatact gaaaatccaa taatcagaaa      420
agtaattttg ccacattatt tatnaaaaat gttcc                                     455

```

```

<210> 721
<211> 530
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(530)
<223> n = A,T,C or G

```

```

<400> 721
ccagtgcctt ctgccgtggt ttagtgattg ggtgtagaa ataaaaactc aggtctatatt      60
cttaccagtc agtaacaatt tttagagaat gtacttggtg tataatatat ggacttcagg      120
aactttattg gggngggggg ttaattttgc cttaccctgt tcactttcag atgattaggc      180
ttttgcactt tagaatgaga aacttgtagc gttagtgtgt tcttactagc ttttaatttgt      240
atgtagcaat gaattgtgaa tcttagtgca gtgggttttt ttaaaaaact caaaaagctg      300
ggaattaagt ggtttcagta ataatgctat accgaggtgc ttgcattgta tttcataatt      360
ttgttacaaa ccaaaattat ttttaatgan aacggtcctg ggttcagagg tgtgatgccg      420
gaatgtatatt tctgactgtt aggcccttgg aacagatacc ggtgctttct tgaaagatga      480
aagaaatgca atgggtgctc ttcatgcaag gttgcaaacc taccaagaat                    530

```

```

<210> 722
<211> 242
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(242)
<223> n = A,T,C or G

```

```

<400> 722
ccaaggggtca tgatggcagg agtaatcana ggtgntcctg tgttggtgata agggngggaga      60
ggttaaagga gccacttatt agtaatgttg atagtagaat gatggctagg gtgacttcat      120
atgagattgt ttgggctact gctcgtagtg cgccgatcag ggcgtagttt gagtttgatg      180
ctcatcctga tnagaggatt gagtaaacgg ctaggctaga ggtggctaga ataaatagga      240
gg                                                    242

```

```

<210> 723
<211> 472
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(472)
<223> n = A,T,C or G

```



```
<210> 724
<211> 292
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(292)
<223> n = A,T,C or G
```

```
<210> 725
<211> 122
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(122)
<223> n = A,T,C or G
```

```
<210> 726
<211> 477
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(477)
<223> n = A,T,C or G
```

<400> 726
ctgaacccctc gtggagccat tcatacaggt ccctaattaa ggaacaagtq attatqctac 60


```
<210> 730
<211> 310
<212> DNA
<213> Homo sapien
```

```
<210> 731
<211> 467
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(467)
<223> n = A,T,C or G
```

```
<210> 732
<211> 492
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(492)
<223> n = A,T,C or G
```

<400> 732
cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaqa 60

```
<210> 733
<211> 562
<212> DNA
<213> Homo sapien
```

| <400> | 733 | | | | | | |
|-------|-------------|------------|------------|------------|-------------|--|-----|
| atggc | aatagcattc | actgtcgtat | tttgcagtg | tcaggaagt | ggacgttaac | | 60 |
| aggtg | cttgtttgta | ttagctctgc | taggtttacc | tctacaacgt | agatttcagc | | 120 |
| cgctg | actgacacta | cattctagtt | cttaagattt | tttttccana | ttcccccttc | | 180 |
| ctaga | catacgtagc | atactttcat | cttattcagt | ctttctgtaa | cctgctgctg | | 240 |
| agtcc | tcctcacctc | agatcggaat | caatggagtg | ggcccagagg | atacatttta | | 300 |
| agtaa | tggtaggtag | atttgtcctg | ctttctaaaa | catctcctca | tttcatattt | | 360 |
| ccata | ttgattccat | aagggaatat | taatgggtgn | ttcctccttt | agggaggcaa | | 420 |
| agagn | gtggacatct | tctaattctg | aggaacagtn | gttgatttcc | cttggaaggag | | 480 |
| atatt | gactgtnttt | ccaataaacc | tgnttgcccc | agntcaatcc | ctcattttaa | | 540 |
| aatgt | tggtntctggg | ct | | | | | 562 |

```
<220>  
<221> misc_feature  
<222> (1)...(265)  
<223> n = A,T,C or G
```

```
<210> 735
<211> 216
<212> DNA
<213> Homo sapien
```

<400> 735

atttaatacg tgctcactgc tcggcacgcg ctgaagctac agttaacaat cagtgagcac 60
 atattaaatg ataaaaataat gctgatggta aacattcata acagcagagt aagattttgg 120
 cagttttgtg tctcggtaac ataactgtaa ccttagatga acacctatcc cttcatgac 180
 tgacttttaga ggcaaggagt ttgtaacatc taatgg 216

<210> 736

<211> 285

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(285)

<223> n = A,T,C or G

<400> 736

ctgaaaggca acntggagac tagttagtct agtccctca tattataaat tggatatgctg 60
 aggccaggca gtaaatgtct atggagctct ccaatttaag gccagtttga ctccaagggt 120
 agggcttcta gtaaaatttt gtgattaaat tggaaaactct aatttatattt tctatgngtt 180
 tttggtacct aatcctcata agcaagccat atttcaaggc tgatcaatga aaacacccaaa 240
 taccaaagct tcctttccct tccaaattta ctgacccttt gtcag 285

<210> 737

<211> 509

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(509)

<223> n = A,T,C or G

<400> 737

agangaagaa gangaagatt aagggaaaag tacatcggtc aagaagagct caacaaaaca 60
 aagcccctct ggaccagaaa tcccgcgat attactaatg aggagtaagg agaattctat 120
 aagagcttga ccaatgactg ggaagatcac ttggcagtga agcatttttc agttgaagga 180
 cagttggaat tcagagccct tctatttgtc ccacgcagtg ctccctttga tctgtttgaa 240
 aacagaaaga aaaagaacaa catcaaattg tatgtacgca gagttttcat catggataac 300
 tngaggagc taatccctga atatctgaac ttcattagag gggtggnaga ctggaggat 360
 ctccctctaa acatatcccg tgagatgttg caacaaagca aaattttgaa agttatcang 420
 aagaatttgg gtcaaaaaat gcttanaact ctttactgaa ctggcggaag atnaagagaa 480
 ctncagana ttctatgagc agntctctt 509

<210> 738

<211> 97

<212> DNA

<213> Homo sapien

<400> 738

cagtgaattg aatacgactc ctatagggcg aattgggcc tctagatgca tgctogagcg 60
 gccgccagtg tgatggatat ctgcagaatt cgccctt 97

<210> 739

<211> 209


```
<210> 743
<211> 275
<212> DNA
<213> Homo sapien
```

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| <400> 743 | | | | | | |
| cangacgcta | cttcccttat | catagaagag | cttatcacct | ttcatgatca | cgccctcata | 60 |
| gtcaattttcc | ttatctgctc | cctagtcctg | tatgcccttt | tcctaacact | cacaacaaaa | 120 |
| ctaactaata | ctaacatctc | agacgctcag | gaaatagaaa | ccgtctgaac | tatcctgccc | 180 |
| gccatcatcc | tagtctcat | cgcctcccca | tccttacgca | tcctttacat | aacagacgag | 240 |
| gtcaacgata | cctcccttac | catcaaata | attgg | | | 275 |

```
<220>  
<221> misc_feature  
<222> (1)...(295)  
<223> n = A,T,C or G
```

```
<210> 745
<211> 477
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(477)
<223> n = A,T,C or G
```

<400> 745
 cgcgttactg tacatattgc tagcaggaga caactggaaa tactaaacaa atactggaat 60
 tcacattaca gacagacgaa accaacatgg atgccacaca taacttcctt tgtagtttca 120
 cagagagcct atttgtggtt gctcagggtg ggtcatacat tgcttgcaga aatggcctga 180
 tcatagctct atgaaacaat gaattcggaa tgaaatctta ccatgacacc tctctgtagg 240
 aaagaaatgt tgcttcacgt gtgctaagtt gagataataa tatttcacat atttatatac 300
 agagaatcac tctcaaattt aacccaagat aagcaatagg atttgggggt gacttgtaga 360
 catttctaac aacacttttc ttttttctag aggtcactct caaactga tatatcacta 420
 tagtttgagt gtanggattc agtaatcaaa ggttggttatt gcaaaagagc caggcag 477

<210> 746
 <211> 524
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(524)
 <223> n = A,T,C or G

<400> 746
 ctgtgaaatt ggggtgggag agccaaaata ctttacaact tcagaccgga gaaaaggcca 60
 gaggtgtgaa gttagactct atgatgaaac agagtctgtt tttgcgatga catggtggga 120
 taatgaatcc attctacttg cacagagctg gatgccacga gaaacagtaa tatttgcctc 180
 agatgtaaga ataaattttg acaaatttcg gaactgcatg acagcaactg taatctcaaa 240
 aaccattatt acaactaatc cagatatacc agaagctaac attctgctga attttatacg 300
 agaaaaataa gaaacaaatg ttctggatga tgaaattgac agttatttca aagaatccat 360
 aaatttaagt acaatagttg atgtctacac agntgaacaa ttaaaggga aagctttgaa 420
 gaatgaagga aaagctgatc cttcctatgg catcctttat gctacattt ccacactcaa 480
 cattgatgat gaaactcaaa agtagttcga aatagatggt ccag 524

<210> 747
 <211> 456
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(456)
 <223> n = A,T,C or G

<400> 747
 cctcagttct tgattgtggt tgacggggcg tcaccatgaa ggagcccatt tagtataaag 60
 cttccaacct tttctcttaa tcgtttcttt aatcttttaa accatcttca agtgcataag 120
 ggagtttccg atgccagagg atgaaagcaa gtgctttctc caccctctcc tcccagagtg 180
 aaaacaaatc cttttgctga tacttgtttc aaaagcatcc attgtaaagc ttctcagtga 240
 cacaaaatac tgagaggtaa ctttttatca atcaaacac ataccccaat ttaacacctt 300
 tcagtgtctc gaattcaact gacagactaa aggggtgttc ctgtaacagt ctgaaatatt 360
 aagtgttttt tttgttttgt ttttaaactc ttttccagaa aacttcctct nggggtagga 420
 aagtacacat gaagcagcaa agtaacgaag aaaaac 456

<210> 748
 <211> 474

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(474)
<223> n = A,T,C or G

<400> 748
ccanaccagg gaaccaaaatg cagacagnga agttctctgc ttcttttggc tataatgnga 60
caagaaaggg atcatctttt gaagatgttt aaagaaataa agcaactttc ttataaaaca 120
gtcaaataat caattaatgg aataaataag tactaaccca cattttaacc actctgtaat 180
cactacactt tacatatattt ttatttnggn ggcaantcc ccataatta gtctaaaatc 240
caccaatcac ttttaaaagt aaaatgaata gccaccaaaa taagaaaatc ttctgttcac 300
tctttggcta aaaaggaaaa caaataaaac aaaacaaaaa gaaacagaag acaactgtaa 360
cactgggtgat aaaagaaact ttttttttac aagtaaaata aagttatcaa tttaaatcct 420
ggncacttta taaaaacaag aggtaatgtt gtaataaaac agcagtagcc tcag 474

<210> 749
<211> 355
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(355)
<223> n = A,T,C or G

<400> 749
cctgggttna gnggctgact gnaacctcca cttcctgttc tcaggcaatc ctctgcctc 60
agcctcetta gtagctggga ctacaggagt gtgcaaccat gcccaactaa tttttgtatt 120
tttaatatag acagggtttc accatgttga tcagggttgt ctccaactcc tgacctcagg 180
tgatccacct gtcccagcct cccaaagtgc tgggattaca ggcatgagcc accacgcccg 240
gnccaggata aagtaaaaaa ttgtaagcac acaaggccct ttgcaacctg gctcctgggt 300
actactttta ncctcctgcc ctcccaaatg tinctactgt ttttctanac atacc 355

<210> 750
<211> 493
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(493)
<223> n = A,T,C or G

<400> 750
ccatgtctggt ctgcaactcc tgaactcagg tgatccaccc gcctcagtct cccaatagat 60
tacatatatt attaatgaat tgcttccttt aacaccctat tcattgaatt ttccagtaaa 120
ccacaattac taattactcc tgaatcaga aaagagggtta aaaagatttt ataacagtat 180
cctatgaaat ctactacttt caagtaatag tagttgaatt accaaaaacc gtcactcaag 240
ccaatgacta caattaagat atgagtaaca tttcctagat aaataaaagtc aattaattat 300
atgtgcatct gggaaataga gaaagtacat ataagccatg attttgaagn caaaagagag 360
agantatttg ccaaggaggg gtgagttata gtatgtaatt ataacataca gaagcttttt 420

```
<210> 751
<211> 364
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|-------|-------------|-------------|------------|------------|------------|--|-----|
| <400> | 751 | | | | | | |
| tctgg | naaggtcacc | aagtctgccc | aganagctca | gaaggctaaa | tgaatattat | | 60 |
| atacc | tgcaccccca | ctcttaatca | gtggtggaag | aacggtctca | gaactgtttg | | 120 |
| attgg | ccatttaagt | ttagtagtaa | aagactgggt | aatgataaca | atgcatcgta | | 180 |
| ttcag | aaggaaaagga | gaatgttttg | nggaccactt | tggttttctt | ttttgcgtgt | | 240 |
| tttta | agttattagt | ttttaaaatc | agtacttttt | aatggaaaca | acttgaccaa | | 300 |
| tgtca | cagaattttg | agaccocatta | aaaaagttaa | atgagataaa | aaaaaaaaan | | 360 |
| | | | | | | | 364 |

```
<220>
<221> misc_feature
<222> (1)...(498)
<223> n = A,T,C or G
```

```
<210> 753
<211> 467
<212> DNA
<213> Homo sapien
```

<400> 753

```

nacaacctta gccanaacca tttacccaaa taaagggata ggcgatagaa attgaaacct      60
ggcgcaatag atatagnacc gcaagggaaa gatgaaaaat tataaccaag cataatatag      120
caaggactaa cccctatacc ttctgcataa tgaattaact agaaataact ttgcaaggag      180
agccaaaagct aagacccccg aaaccagacg agctatctaa gaacagctaa aagagcacac      240
ccgtctatgt agcaaaatag tgggaagatt tataggtaga ggcgacaaac ctaccgagcc      300
tgggtgatagc tggntgncca agatagaatc ttagntcaac tttaaatttg cccacagaac      360
cctctaaatc cccttgtaaa ttttaactgtt agtccaaaga ggaacagctc ttggacacna      420
ggaaaaaacc ttgcagagag agtaaaaaat ttaacacca tagtagg                      467

```

```

<210> 754
<211> 196
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(196)
<223> n = A,T,C or G

```

```

<400> 754
gtcatgttca agtgttntaa tctgacgcag gcttatgcgg aggagaatgt tttcatgtta      60
cttatactaa cattagttct tctatagggt gatagattgg tccaattggg tgtgaggagt      120
tcagttatat gtttgggatt ttttaggcag tgggtggtga gcttgaacgc tttcttaatt      180
ggtggctgct tttagg                      196

```

```

<210> 755
<211> 381
<212> DNA
<213> Homo sapien

```

```

<400> 755
ctggaaagga ttctgtacat ataagacatc aaatattgag ggatactgga actttttaat      60
taatgggcaa agaaagtcaa caaaggaagt tcatatgaaa tcaaactagt aatatgatta      120
caaaaaaaaaa gtttaaaatt tttcttggcc ccagtcttat catttctgag ccaaatacaa      180
ttctatcgaa atcacctgaa actgaaatca ccattctagg ctgggttttc cataaagatg      240
gactgctcca aaaagaggaa tcaagaaaga atttggctca cagtgaatta ttcaacttgt      300
cttagttaac taaaaataaa atctgactgt taactacaga aatcatttca aattctgtgg      360
tgataataaa gtaatgaccg c                      381

```

```

<210> 756
<211> 341
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(341)
<223> n = A,T,C or G

```

```

<400> 756
ggntataaac ctattattta ttgcagaact aataaaaaat ccaaagcctt gtatttgtac      60
atctttatta tctctaaagc actttcctca acctaatttc agtttttaca attggtactc      120
aagaaaaatag agacagaaat catttgattt tgcccagaaa ccatctgctt atatttataa      180
ggccacctaa tttgaaatca catatagacc aggcgcggtg gtcacgcct gtaattocaa      240

```

```
<210> 757
<211> 479
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(479)
<223> n = A,T,C or G
```

```
<210> 758
<211> 267
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(267)
<223> n = A,T,C or G
```

```
<210> 759
<211> 449
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(449)
<223> n = A,T,C or G
```

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| <400> | 759 | | | | | |
| cgaggtcttg | aaatcagcaa | cacacttaca | aatgagaaaa | tgaaaataga | agagtatata | 60 |
| aagaaaagga | aagaggatta | tgaagagagt | catcagagag | ctgtggctgc | agaggatatcc | 120 |
| gtacttgaaa | actggaagga | gagtgaagtg | tataagctac | agatcatgga | gtcacaaagca | 180 |

```
<210> 760
<211> 414
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(414)
<223> n = A,T,C or G
```

```
<210> 761
<211> 428
<212> DNA
<213> Homo sapien
```

```
<210> 762
<211> 574
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(574)
<223> n = A,T,C or G
```

<400> 762

| | | | | | | |
|-------------|------------|------------|------------|-------------|------------|-----|
| caggctctgaa | ctgataagta | ttaagagacg | tttgttgcta | gttaaagngtt | ccagttgaga | 60 |
| gttcgaagtg | aaaacctggg | ctctttacca | gtgttgagtg | agaagattta | tttctctttc | 120 |
| ctctgaattt | accacatgta | acatcacaga | gacatgtaga | gttcctttag | gatttgcat | 180 |

```

ttgaaccagn ccagtctgat tttcaggtga attctgtgaa gagcttgatg ggggaagtct 240
gaagacagaa ggaattaggg aaaaggggtga tacttacaga gtaaaggaaa taaatgaaaa 300
gataatggta tttttggtag ccacagggaa atagcaggag gggactggag atcacacaca 360
cgcacacgca cacacacaaa cacacacaca cgctaaaact caaactaaaa acctcccaa 420
ggagctgctt tgtttgcaga cttcaattng aagtagatac taagggaag aatagaccag 480
ttaaattca cctgaaaatc tcttcccann cttcaaatgt gctaaaatat cactgtcagc 540
ttagcatctc tncatgtatg tatatataga tgta 574

```

```

<210> 763
<211> 465
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(465)
<223> n = A,T,C or G

```

```

<400> 763
cctactatgg gtgttaaaat tttttactct ctctacaagg ntttttcta gtgtccaaag 60
agctgttctt ctttggaacta acagttaaat ttacaagggg atttagaggg ttctgnngggc 120
aaatttaaaag ttgaactaag attctatctt ggacaaccag ctatcaccag gctcggtagg 180
tttgctgcct ctacctataa atcttcccac tattttgcta catagacggg tgtgctcttt 240
tagctgttct taggtagctc gtctggtttc ggggggtctta gctttggctc tccttgcaaa 300
gttatttcta gttaattcat tatgcagaag gtataggggt tagtccttgc tatattatgc 360
ttggatataa tttttcatct ttcccttgcg gtactatatac tattgcgcca ngtttcaatt 420
tctatcgctt atactttatt tgggtaaatg gtttggtctaa ggttg 465

```

```

<210> 764
<211> 151
<212> DNA
<213> Homo sapien

```

```

<400> 764
ctgtcaatta atgctagtcc tcaggattta aaaaataatc ttaactcaaa gtccaatgca 60
aaaacattaa gttggtaatt actcttgatc ttgaattact tccggtacga aagtccttca 120
catttttcaa actaagctac tatatttaag g 151

```

```

<210> 765
<211> 251
<212> DNA
<213> Homo sapien

```

```

<400> 765
gaagagctta tcacctttca tgatcacgcc ctcatagtca ttttccttat ctgcttcta 60
gtcctgtatg cctttttcct aacactcaca acaaaaactaa ctaataactaa catctcagac 120
gctcaggaaa tagtaaccgt ctgaactatc ctgcccgcga tcatcctagt cctcatcgcc 180
ctcccatccc tacgcatcct ttacataaca gacgaggtca acgatccctc ccttaccatc 240
aatcaattg g 251

```

```

<210> 766
<211> 375
<212> DNA
<213> Homo sapien

```

<220>
 <221> misc_feature
 <222> (1)...(375)
 <223> n = A,T,C or G

<400> 766
 cgaggtctgn cctcctgggt cttcatccat tattaacaga agagcatact ggtttcgggc 60
 cataaaatct ttgggaaggg acaactgtaa aggaagttca tagtcgtcaa tatgaaggat 120
 ttttaatttct ggctttccta tcttcttctt caggatagct tccttcagca tagaattggt 180
 ttccaatata aaatattttg ctgggttggt cgtactatgt aggctgacca ctgggaccct 240
 tggaccttca cagaataata agaaatgttg attcatggga ctaaaactgg catcaaaata 300
 tgtacattgt tctttcatga aattacatga aatgcattgg cgattcaata atccttcagt 360
 agaagcactg tacag 375

<210> 767
 <211> 485
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(485)
 <223> n = A,T,C or G

<400> 767
 cgaggtctga accctcgtgg agccattcat acaggtccct aattaaggaa caagtgatta 60
 tgctaccttn gcacgggttag ggtaccgcgg cccgttaaac atgtgtcact gggcaggcgg 120
 tgcctctaact actggtgatg ctagaggtga tgtttttggn aaacaggcgg ggtaagattt 180
 gccgagttcc ttttactttt ttttaacctt ctttatgagc atgcctgtgt tgggttgaca 240
 gtgagggtaa taatgacttg ttggtgattg tagatatggt gctgttaatt gtcagttcag 300
 tgttttaate tgacgcaggc ttatgcggag gagaatgttt tcatgttact tatactaaca 360
 ttagttcttc tatagggtga tagatnggtc caattgggtg tgaggagntc acttatatgt 420
 ttgggatttt ttaggtaagn ggggtgtgag cttgaacgct ttcttaattg ggggctgctt 480
 ttang 485

<210> 768
 <211> 379
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(379)
 <223> n = A,T,C or G

<400> 768
 ctgatattct attaaagata caaagaggag ctggnaccat ttcttctgaa actattacaa 60
 acaactgaaa aggtggaatt tctccctaact tcatttttagg aggccagcat tatactgata 120
 ccaaaacctg gcagaggtac aataataaaa ggaaacttca agtcagtatc actgatgaac 180
 accaatgtga aaatcctcaa taaaatactg gcaaactgaa ttcagcagca catcaaaaag 240
 ctaatccacc acaatcaagt cagcttcac cctgcgatgc aagtctggtt caacatatgc 300
 aaatcaataa atacaattca tcagataaac agagctaaag acaaaattca catgattttc 360
 tcaatagatg cagaaaagg 379

<210> 769
 <211> 518
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(518)
 <223> n = A,T,C or G

<400> 769
 cgagggtccat atgatgatca gtctatatag ttttaaggcgc agatacacaa attttcaaaa 60
 atatgggtag aatatagtca atatgaatgg aatagacaat gctttgaaaa tcaactggagg 120
 gaggccttat tgtttgtgaa aacatgttgt catcactttt tgctttaagc ccttggtggt 180
 gaaataactc aaaccattct tccttatgct gaagatcgag aaccccaagt atcacatcta 240
 ccatcccact catcaatgtg attggtcagt ctttgcagag gncctgcata gccagtttta 300
 aagtttaggt tcttgcata acatatgaaa aggcattgta cttgtgcttt caaagagctt 360
 tttgcttggg gtaaaaagaa aactcaaatt acagtgtgat gtggaatata atgggtgtag 420
 tttcatcgag atgatgggaa agaattgata agataaagcn gaaagatgag cagaattttc 480
 agattgggtn tggaaagagc acttaagaaa gaggggtgg 518

<210> 770
 <211> 378
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(378)
 <223> n = A,T,C or G

<400> 770
 tatgggtcct gagtgtggaa tataagataa caagacaatt cccttgcttt caagggaaat 60
 cacactttat aaaactttga attcctgaaa tgggtttcag aggttccaag gtcaaattca 120
 agaataagag ttaagaagaa aaagactatg agaaaggaag tgntgacccc atttgcattt 180
 aaatggcagg aatagtctca atctactcat tggggaaaaa tgtatgttgc atatttttga 240
 gatattgcaa cttgctctct ctctttgcca ccccaccctt tgncatgctc tgtttttggg 300
 ctgaattggc aagaaaaatg gctggagggc tggagaaggn tggacccttc ttccttcttc 360
 cttcttcctt ctttctcc 378

<210> 771
 <211> 207
 <212> DNA
 <213> Homo sapien

<400> 771
 cataaatatt atactagcat ttaccatctc acttctagga atactagtat atcgctcaca 60
 cctcataatcc tccttactat gcctagaagg aataatacta tcaactgttca ttatagctac 120
 tctcataacc ctcaacaccc actcctctct agccaatatt gtgcctattg ccatactagt 180
 ctttgcogcc tgcgaagcag cggtagg 207

<210> 772
 <211> 384

<400> 772

<400> 773

<400> 774

<400> 775

```
ccatggctaa gntatataga tagctgggtg gctggagtaa atgantgagg nacgagtcctg      60
angagggttag ttgaggcaat aaaaatgatn aaggatacta gtataagaga tcangttcgt      120
cctttacatg ttgngtatgg ctatcatttg ttttgaggct agnttgatta gtcattgttg      180
ggtggttaatt aa                                     192
```

| | |
|-------|-----|
| <210> | 779 |
| <211> | 277 |
| <212> | DNA |

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(277)

<223> n = A,T,C or G

<400> 779

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctnttgatt | tgatgggtaa | ggggagggat | cgttgacctc | gtctggtatg | taaaggatgc | 60 |
| gtagggatgg | gagggcgatg | aggactagga | tgatggcggg | caggatagtt | cagacggttt | 120 |
| ctatttcctg | agcgtctgag | atgttagtat | tagttagttt | tgttgtgagt | gttaggaaaa | 180 |
| gggcatacag | gactaggaag | cagataagga | aaatgactat | gagggcgtga | tcatgaaagg | 240 |
| tgataagctc | ttctatgata | ggggaagtag | cgtcttg | | | 277 |

<210> 780

<211> 328

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(328)

<223> n = A,T,C or G

<400> 780

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| catgntatgg | ataacatnt | taactgtatt | ttntgcanc | cgtaccttct | tgggaataca | 60 |
| attgtctaac | tttttat | ttt | tggnctggct | ggttggtgt | gcaaaactcc | 120 |
| atgttgccac | actgcaacac | cttacagatg | tggaagatgt | gaaatttgtc | atcaattatg | 180 |
| actaccctaa | ctcctcagag | gattatattc | atcgaattgg | aagaactgct | cgcagtacca | 240 |
| aaacaggcac | agcatacact | ttctttacac | ctaataacat | aaagcagggg | agcgacctta | 300 |
| tctctgtgct | tcgggaagct | aancaaac | | | | 328 |

<210> 781

<211> 305

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(305)

<223> n = A,T,C or G

<400> 781

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgttcagaa | agctcattgg | acctggtttt | gaaaataaaa | caaagttaaa | accctgggag | 60 |
| gagttattgt | gcagngtgga | gtactcaggc | tttcttataa | agaaaaaaaa | agttatctgg | 120 |
| taccaaagtg | tgcaacctac | agaccctcag | gtactgccct | gtgacttctc | tgtatgacat | 180 |
| cacaaggctg | ccaagtgcct | gtttttctag | aactaggagt | tggtgagggt | tggctantgc | 240 |
| tgaaaccatg | cataggattg | gtttactaaa | ttaaaacctt | attacgtacg | tcctccaaaa | 300 |
| gacag | | | | | | 305 |

<210> 782

<211> 497

<212> DNA

<213> Homo sapien

005250 E3T3E0

<220>
 <221> misc_feature
 <222> (1)...(497)
 <223> n = A,T,C or G

<400> 782
 cgaggtggct ttaattgatg ttaatgcctt atgtcaaattg taaagttaga atttgctagg 60
 gctgggatag ggagtgatat ttctaggact tagacattga aaactaattc agcctgtagt 120
 aacctggatg gttttcaatg gcatggttag tcaaattcat ggttttaaac ttagaagcag 180
 ctttcggggg agagggtagg ttggagcatt tattacatat tttactgttt aatgtcttaa 240
 ccgtgggcct ttttaatttgt aaacactgaa atgattgttg ggctgtggaa aacatttacc 300
 tatttacctt ggaagtttta aaagacagtc cacttttttag catgtgtgtt gcgtccagcc 360
 tgtggtcgtc ttaactaata aatgngattt ttctctcaa aaaaaaacct ccccgggcgg 420
 ccgctcaagg gcaaatccn cacactggcg gccgttacta ggggatccga nctcgggtcca 480
 agcttggcgt aatcatg 497

<210> 783
 <211> 364
 <212> PRT
 <213> Homo sapien

<400> 783
 Met Trp Gln Pro Leu Phe Phe Lys Trp Leu Leu Ser Cys Cys Pro Gly
 1 5 10 15
 Ser Ser Gln Ile Ala Ala Ala Ala Ser Thr Gln Pro Glu Asp Asp Ile
 20 25 30
 Asn Thr Gln Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val Thr Asp
 35 40 45
 Ser Pro Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr Ser Ser His
 50 55 60
 Gly Ala Asn Arg Phe Val Pro Lys Ser Lys Ala Leu Glu Ala Val Lys
 65 70 75 80
 Leu Ala Ile Glu Ala Gly Phe His His Ile Asp Ser Ala His Val Tyr
 85 90 95
 Asn Asn Glu Glu Gln Val Gly Leu Ala Ile Arg Ser Lys Ile Ala Asp
 100 105 110
 Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
 115 120 125
 Asn Ser His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Arg Ser Leu
 130 135 140
 Lys Asn Leu Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Phe Pro
 145 150 155 160
 Val Ser Val Lys Pro Gly Glu Glu Val Ile Pro Lys Asp Glu Asn Gly
 165 170 175
 Lys Ile Leu Phe Asp Thr Val Asp Leu Cys Ala Thr Trp Glu Ala Met
 180 185 190
 Glu Lys Cys Lys Asp Ala Gly Leu Ala Lys Ser Ile Gly Val Ser Asn
 195 200 205
 Phe Asn His Arg Leu Leu Glu Met Ile Leu Asn Lys Pro Gly Leu Lys
 210 215 220
 Tyr Lys Pro Val Cys Asn Gln Val Glu Cys His Pro Tyr Phe Asn Gln
 225 230 235 240
 Arg Lys Leu Leu Asp Phe Cys Lys Ser Lys Asp Ile Val Leu Val Ala

```
<210> 784
<211> 6353
<212> DNA
<213> Homo sapien
```

| | | | | | | |
|------------|-------------|------------|-------------|------------|-------------|------|
| <400> 784 | | | | | | |
| tggcgaatgg | gacgcgccct | gtagcggcgc | attaagcgcg | gcgggtgtgg | tggttacgcg | 60 |
| cagcgtgacc | gctacacttg | ccagcgcctt | agcgcccgct | cctttcgctt | tcttcccttc | 120 |
| ctttctcgcc | acgttcgcgc | gctttccccg | tcaagctcta | aatcgggggc | tcccttttagg | 180 |
| gttccgattt | agtgtctttac | ggcacctcga | ccccaaaaaa | cttgattagg | gtgatggttc | 240 |
| acgtagtggg | ccatcgccct | gatagacggg | ttttcgccct | ttgacgttgg | agtccacggt | 300 |
| ctttaatatg | ggactcttgt | tccaaactgg | aacaacactc | aaccctatct | cgggtctattc | 360 |
| ttttgattta | taagggtatt | tgccgatttc | ggcctatttg | ttaaaaaatg | agctgattta | 420 |
| acaaaaattt | aacgcgaatt | ttaacaaaat | attaacgttt | acaatttcag | gtggcacttt | 480 |
| tccgggaaat | gtgcgcggaa | cccctatttg | tttatttttc | taaatacatt | caaatatgta | 540 |
| tccgctcatg | aattaattct | tagaaaaact | catcgagcat | caaatgaaac | tgcgaatttat | 600 |
| tcatatcagg | attatcaata | ccatattttt | gaaaaagccg | tttctgtaat | gaaggagaaa | 660 |
| actcaccgag | gcagttccat | aggatggcaa | gaccttggtg | tccgtctgcg | attccgactc | 720 |
| gtccaacatc | aatacaacct | attaattttc | cctcgctcaa | aataaggtta | tcaagtgaga | 780 |
| aatcaccatg | agtgcgcact | gaatccggtg | agaatggcaa | aagtttatgc | atttcttttc | 840 |
| agacttgttc | aacaggccag | ccattacgct | cgtcatcaaa | atcactcgca | tcaaccaaac | 900 |
| cgttattcat | tctgtattgc | gcctgagcga | gacgaaatac | gcgatcgctg | ttaaaggagc | 960 |
| aattacaaac | aggaatcgaa | tgcaaccggc | gcaggaacac | tgccagcgca | tcaacaatat | 1020 |
| tttcacctga | atcaggatat | tcttctaata | cctggaatgc | tgttttcccg | gggatcgcg | 1080 |
| tggtgagtaa | ccatgcacga | tcaggagtag | ggataaaatg | cttgatggtc | ggaagaggca | 1140 |
| taaattccgt | cagccagttt | agtctgacca | tctcatctgt | aacatcattg | gcaacgctac | 1200 |
| ctttgccatg | tttcagaaac | aactctggcg | catcgggctt | cccatacaat | cgatagattg | 1260 |
| tgcacactga | ttgcccgcga | ttatcgcgag | cccattttata | cccataataa | tcagcatcca | 1320 |
| tgttggaatt | taatcgcggc | ctagagcaag | acgtttcccg | ttgaatatgg | ctcataacac | 1380 |
| cccttgattt | actgtttatg | taagcagaca | gttttattgt | tcatgaccac | aatcccttaa | 1440 |
| cgtgagtttt | cgttccactg | agcgtcagac | cccgtagaaa | agatcaaagg | atcttcttga | 1500 |
| gacccctttt | ttctgcgcgt | aatctgctgc | ttgcaaacaa | aaaaaccacc | gctaccagcg | 1560 |
| gtggtttgtt | tgccggatca | agagctacca | actctttttc | cgaaggtaac | tggcttcagc | 1620 |
| agagcgcaga | taccaaatac | tgtccttcta | gtgtagccgt | agttaggcca | cgaactcaag | 1680 |
| aaactctgtg | caccgcctac | atacctcgct | ctgctaattc | tgttaccagt | ggctgctgcc | 1740 |
| agtggcgata | agtcgtgtct | taccgggttg | gactcaagac | gatagttacc | ggataaggcg | 1800 |
| cagcggtcgg | gctgaacggg | gggttcgtgc | acacagccca | gcttgagcgc | aacgacctac | 1860 |
| accgaactga | gatacctaca | gcgtgagcta | tgagaaagcg | ccacgcttcc | cgaagggaga | 1920 |

| | | | | | | |
|-------------|------------|------------|-------------|------------|-------------|------|
| aaggcggaca | ggtatccggt | aagcggcagg | gtcggaaacag | gagagcgcac | gagggagctt | 1980 |
| ccagggggaa | acgcctggta | tctttatagt | cctgtcgggt | ttcgccacct | ctgacttgag | 2040 |
| cgtcgatttt | tgtgatgctc | gtcagggggg | cggagcctat | ggaaaaacgc | cagcaacgcg | 2100 |
| gcctttttac | ggttcctggc | cttttgctgg | ccttttgctc | acatgttctt | tctgcgtta | 2160 |
| tcccttgatt | ctgtggataa | ccgtattacc | gcctttgagt | gagctgatac | cgctcgccgc | 2220 |
| agccgaacga | ccgagcgcag | cgagtcagtg | agcgaggaag | cggaagagcg | cctgatgcgg | 2280 |
| tattttctcc | ttacgcacat | gtgcggtatt | tcacaccgca | tatatggtgc | actctcagta | 2340 |
| caatctgctc | tgatgccgca | tagttaagcc | agtatacact | ccgctatcgc | tacgtgactg | 2400 |
| ggatcatggct | gcgccccgac | acccgccaac | acccgctgac | gcgccccgac | gggcttgtct | 2460 |
| gctccccgga | tccgcttaca | gacaagctgt | gaccgtctcc | gggagctgca | tgtgtcagag | 2520 |
| gttttcaccg | tcacaccga | aacgcgcgag | gcagctgcgg | taaagctcat | cagcgtggtc | 2580 |
| gtgaagcgat | tcacagatgt | ctgcctgttc | atccgcgtcc | agctcgttga | gtttctccag | 2640 |
| aagcgtaaat | gtctggcttc | tgataaagcg | ggccatgtta | agggcggttt | tttctgttt | 2700 |
| ggctactgat | gcctccgtgt | aagggggatt | tctgttcatg | ggggtaatga | taccgatgaa | 2760 |
| acgagagagg | atgctcacga | tacgggttac | tgatgatgaa | catgcccggt | tactggaacg | 2820 |
| ttgtgagggt | aaacaactgg | cggtatggat | gcggcgggac | cagagaaaaa | tactcaggg | 2880 |
| tcaatgccag | cgcttcgtta | atacagatgt | aggtgttcca | cagggtagcc | agcagcatcc | 2940 |
| tgcatgacag | atccggaaca | taatggtgca | ggcgctgac | ttccgcgttt | ccagacttta | 3000 |
| cgaaacacgg | aaaccgaaga | ccattcatgt | tgttgctcag | gtcgcagacg | ttttgcagca | 3060 |
| gcagtcgctt | cacgttcgct | cgctatcgg | tgattcattc | tgctaaccag | taaggcaacc | 3120 |
| ccgccagcct | agccgggtcc | tcaacgacag | gagcacgac | atgcgcaccc | gtggggccgc | 3180 |
| catgccggcg | ataatggcct | gcttctcgcc | gaaacgtttg | gtggcgggac | cagtgcgaa | 3240 |
| ggcttgagcg | agggcggtga | agattccgaa | taccgcaagc | gacaggccga | tcacgtcgcc | 3300 |
| gctccagcga | aagcgttcct | cgccgaaaat | gaccagagc | gctgcgggca | cctgtcctac | 3360 |
| gagttgcatg | ataaagaaga | cagtcataag | tgccgagacg | atagtcatgc | cccgccgcca | 3420 |
| ccggaaggag | ctgactgggt | tgaaggctct | caagggcac | ggtcagatc | ccggtgecta | 3480 |
| atgagtgagc | taacttacat | taattgcgtt | gcgctcactg | cccgtttcc | agtcgggaaa | 3540 |
| cctgtcgtgc | cagctgcatt | aatgaatcgg | ccaacgcgcg | gggagaggcg | gtttgcgtat | 3600 |
| tggggccag | ggtggttttt | cttttcacca | gtgagacggg | caacagctga | ttgcccttca | 3660 |
| ccgcctggcc | ctgagagagt | tgacgcaagc | ggtccacgct | ggtttgcccc | agcaggcgaa | 3720 |
| aatcctgttt | gatggtggtt | aacggcggga | tataacatga | gctgtcttcg | gtatcgctgt | 3780 |
| atcccactac | cgagatatcc | gcaccaacgc | gcagcccga | ctcggtaatg | gcgcgcattg | 3840 |
| cgcccagcgc | catctgatcg | ttggcaacca | gcacgcgagt | gggaacgatg | ccctcattca | 3900 |
| gcattttgat | ggttttgttg | aaaccggaca | tggcactcca | gtcgccttcc | cgttccgcta | 3960 |
| tcggctgaat | ttgattgcga | gtgagatatt | tatgccagcc | agccagacgc | agacgcgcgc | 4020 |
| agacagaact | taatgggccc | gctaacagcg | cgatttgctg | gtgacccaat | gcgaccagat | 4080 |
| gctccacgcc | cagtcgcgta | ccgtcttcat | gggagaaaat | aatactgttg | atgggtgtct | 4140 |
| ggtcagagac | atcaagaaat | aacgcgcgga | cattagtgc | ggcagcttcc | acagcaatgg | 4200 |
| catcctggtc | atccagcgga | tagttaatga | tcagcccact | gacgcgttgc | gcgagaagat | 4260 |
| tgtgcaccgc | cgctttacag | gcttcgacgc | cgcttcgttc | taccatcgac | accaccacgc | 4320 |
| tggcaccag | ttgatcggcg | cgagatttaa | tcgccgcgac | aatttgcgac | ggcgcggtgca | 4380 |
| gggccagact | ggaggtggca | acgcgaatca | gcaacgactg | tttgcccgc | agttgttgtg | 4440 |
| ccacgcgggt | gggaatgtaa | ttcagctccg | ccatcgccgc | ttccactttt | tcccgcgttt | 4500 |
| tcgcagaaac | gtggtggtgc | tggttcacca | cgcgggaaac | ggtctgataa | gagacaccgg | 4560 |
| catactctgc | gacatcgat | aacgttactg | gtttcacatt | caccaccctg | aattgactct | 4620 |
| cttcggggcg | ctatcatgcc | ataccgcgaa | aggttttgcg | ccattcgatg | gtgtccggga | 4680 |
| tctcgacgct | ctcccttatg | cgactcctgc | attaggaagc | agcccagtag | taggttgagg | 4740 |
| ccgttgagca | ccgcgcgcgc | aaggaatggt | catagcgaag | agatggcgcc | caacagtcgc | 4800 |
| ccggccacgg | ggcctgccac | cataccacgc | ccgaaacaag | cgctcatgag | cccgaagtgg | 4860 |
| cgagcccgat | cttcccatc | ggtgatgtcg | gcgatatagg | cgccagcaac | cgcacctgtg | 4920 |
| gcgcgggtga | tgccggccac | gatgcgtccg | gcgtagagga | tcgagatctc | gatcccgcca | 4980 |
| aattaatacg | actcactata | ggggaattgt | gagcggataa | caattccctc | ctagaaataa | 5040 |
| ttttgtttaa | ctttaagaag | gagatataca | tatgcagcat | caccaccatc | accactggca | 5100 |
| gcccctcttc | ttcaagtggc | tcttgtctcg | ttgcctggg | agttctcaaa | ttgctgcagc | 5160 |

```
<210> 785
<211> 5502
<212> DNA
<213> Homo sapien
```

| | | | | | | |
|-------------|--------------|-------------|--------------|-------------|-------------|------|
| <400> 785 | | | | | | |
| tggcgcaatgg | gacgcgcocct | gtagcgggcgc | attaagcgcg | gcgggtgtgg | tggttacgcg | 60 |
| cagcgtgacc | gctacacttg | ccagcgcocct | agcgcocgcct | ccttttcgctt | tcttcccttc | 120 |
| ctttctcgcc | acgttcgcgcg | gcttttccccg | tcaagctcta | aatcggggggc | tcccttttagg | 180 |
| gttccgattt | agtgtctttac | ggcacctcga | ccccaaaaaa | cttgattagg | gtgatggttc | 240 |
| acgtagtggg | ccatcgccocct | gatagacggg | ttttcgccocct | ttgacgttgg | agtccacggt | 300 |
| ctttaatagt | ggactccttg | tccaaactgg | aacaacactc | aaccttatct | cgggtctattc | 360 |
| ttttgattta | taagggtatt | tgcgcgatttc | ggcctattgg | ttaaaaaatg | agctgattta | 420 |
| acaaaaattt | aacgcgaatt | ttaacaaaaat | attaacgttt | acaattttcag | gtggcacttt | 480 |
| tcgggggaaat | gtgcgcggaa | cccctatttg | tttatttttc | taaatacatt | caaataatgta | 540 |
| tcgcgtcatg | aattaattct | tcgaaaaact | catcgagcat | caaatgaaac | tgaacatttat | 600 |
| tcatatcagg | attatcaata | ccatatTTTT | gaaaaagcgc | tttctgtaat | gaaggagaaa | 660 |
| actcaccgag | gcagttccat | aggatggcaa | gacctctggt | tcggtctgcg | attccgactc | 720 |
| gtccaacatc | aatacaacct | attaatttcc | cctcgtcaaa | aataaggtta | tcaagtgaga | 780 |
| aatcaccatg | agtgcgcgact | gaatccgggtg | agaatggcaa | aagtttatgc | atttcttttc | 840 |
| agacttggtc | aacaggccag | ccattacgct | cgctcatcaa | atcactcgca | tcaaccaaac | 900 |
| cgttattcat | tcgtgattgc | gcctgagcga | gacgaaatac | gcgatcgctg | ttaaaggagc | 960 |
| aattacaaac | aggaatcgaa | tgcaaccggc | gcaggaacac | tgccagcgca | tcaacaatat | 1020 |
| tttcacctga | atcaggatat | tcttctaata | cctggaatgc | tgttttccccg | gggatcgcag | 1080 |
| tggtgagtaa | ccatgcatca | tcaggagtac | ggataaaatg | cctgatgggtc | ggaagaggca | 1140 |
| taaattccgt | cagccagttt | agtctgacca | tctcatctgt | aacatcattg | gcaacgctac | 1200 |
| ctttgccatg | tttcagaaac | aactctggcg | catcgggctt | cccatacaat | cgatagattg | 1260 |
| tcgcacctga | ttgcccgaca | ttatcgcgag | cccatttata | cccataaaa | tcagcatcca | 1320 |
| tccttggaaat | taatcgcggc | ctagagcaag | acgtttccccg | ttgaatatgg | ctcataacac | 1380 |
| cgcttgattt | actgtttatg | taagcagaca | gttttattgt | tcatgaccaa | aatcccttaa | 1440 |
| cgtgaagttt | cgttccactg | aqcgtcagac | cccgtagaaa | agatcaaagg | atcttcttga | 1500 |

gatcccttttt ttctgcgcgt aatctgctgc ttgcaaacaa aaaaaccacc gctaccagcg 1560
gtggtttgtt tgccggatca agagctacca actctttttc cgaaggtaac tggcttcagc 1620
agagcgcaga taccaaatac tgtccttcta gtgtagccgt agttaggcca ccacttcaag 1680
aactctgtag caccgcctac atacctcgct ctgctaatac tgttaccagt ggctgctgcc 1740
agtggcgata agtcgtgtct taccgggttg gactcaagac gatagttacc ggataaggcg 1800
cagcggtcgg gctgaacggg gggttcgtgc acacagccca gcttggagcg aacgacctac 1860
accgaactga gatacctaca gcgtgagcta tgagaaagcg ccacgcttcc cgaagggaga 1920
aaggcggaca ggtatccggt aagcggcagg gtccggaacag gagagcgcac gagggagctt 1980
ccagggggaa acgcctggta tctttatagt cctgtcgggt ttcgccacct ctgacttgag 2040
cgtcgatttt tgtgatgctc gtcagggggg cggagcctat ggaaaaacgc cagcaacgcg 2100
gccttttttac ggttcctggc cttttgctgg ccttttgctc acatgttctt tctcgctta 2160
tcccctgatt ctgtggataa ccgtattacc gcctttgagt gagctgatac cgctcgccgc 2220
agccgaacga ccgagcgcag cgagtcagt agcgaggaag cggagagcg cctgatgcgg 2280
tattttctcc ttacgcatct gtgcggtatt tcacaccgca tatatgggtc actctcagta 2340
caatctgtc tgatgccga tagttaagcc agtatacact ccgctatcgc tacgtgactg 2400
ggtcatggct gcgccccgac acccgccaac acccgctgac gcgccccgac gggcttgtct 2460
gctcccggca tccgcttaca gacaagctgt gaccgtctcc gggagctgca tgtgtcagag 2520
gttttcaccg tcataccga aacgcgcgag gcagctgcgg taaagctcat cagcgtggct 2580
gtgaagcgat tcacagatgt ctgcctgttc atccgcgtcc agctcgttga gtttctccag 2640
aagcgttaat gtctggcttc tgataaagcg ggccatgtta agggcggtt tttcctgttt 2700
ggctactgat gctcctgtgt aagggggatt tctgttcatg ggggtaatga taccgatgaa 2760
acgagagagg atgtcacga tacgggttac tgatgatgaa catgcccggg tactggaacg 2820
ttgtgaggg aaacaactgg cggatggat gcggcgggac cagagaaaaa tcaactcagg 2880
tcaatgccag cgcttcgtta atacagatgt aggtgttcca cagggtagcc agcagcatcc 2940
tgcatgagc atccggaaca taatggtgca gggcgctgac ttccgcgttt ccagacttta 3000
cgaaacacgg aaaccgaaga ccattcatgt tgttgcctag gtcgcagacg ttttgacga 3060
gcagtcgctt cagcttcgct cgcgtatcgg tgattcattc tgctaaccag taaggcaacc 3120
ccgccagcct agccgggtcc tcaacgacag gagcacgatc atgcgcaccc gtggggccgc 3180
catgccggcg ataagggcct gcttctcgcc gaaacgtttg gtggcgggac cagtgaacga 3240
ggcttgagcg agggcggtgca agattccgaa taccgcaagc gacaggccga tcatcgtcgc 3300
gctccagcga aagcggctct cgccgaaaa gaccagagc gctgcgggca cctgtcctac 3360
gagttgcatg ataaagaaga cagtcataag tgcggcgacg atagtcatgc ccgcgcccc 3420
ccgaaggag ctgactgggt tgaaggctct caagggcata ggtcgagatc ccggtgccta 3480
atgagtgagc taacttacat taattgcgtt gcgctcactg cccgctttcc agtcgggaaa 3540
cctgtcgtgc cagctgcatt aatgaatcgg ccaacgcgcg gggagaggcg gtttgcgtat 3600
tgggcgccag ggtggttttt cttttacca gtgagacggg caacagctga ttgcccttca 3660
ccgcctggcc ctgagagagt tgcagcaagc ggtccacgct gggttgcccc agcaggcgaa 3720
aatcctgttt gatggtggtt aacggcggga tataacatga gctgtcttcg gtatcgtcgt 3780
atccactac cgagatatcc gcaccaacgc gcagcccgga ctcggtaatg gcgcgcatg 3840
cgcccagcgc catctgatcg ttggcaacca gcacgcagc ggaacgatg cctcattca 3900
gcatttgcat ggtttgttga aaaccggaca tggcactcca gtcgccttcc cgttccgcta 3960
tcggctgaat ttgattgca gtgagatatt tatgccagcc agccagacgc agacgcgcgc 4020
agacagaact taatgggccc gctaacagcg cgatttgctg gtgacccaat gcgaccagat 4080
gtccacgcgc cagtcgcgta ccgtcttcat gggagaaaat aatactgttg atgggtgtct 4140
ggtcagagac atcaagaaat aacgcgggaa cattagtga ggcagcttcc acagcaatgg 4200
catcctggtc atccagcgga tagttaatga tcagccact gacgcgttgc gcgagaagat 4260
tgtgacccgc cgctttacag gcttcgacgc cgcttcgttc taccatcgac accaccgc 4320
tggcaccag ttgatcggcg cgagatttaa tcgcgcgac aattgcgac ggcgcgtgca 4380
ggcccgact ggaggtggca acgccaatca gcaacgactg ttgcccgc agttgttgtg 4440
ccacgcaggt ggggaatgtaa ttcagctccg ccacgcgcgc ttccactttt tccgcgttt 4500
tcgcagaaac gtggctggcc tggttacca cgcgggaaac ggtctgataa gagacaccgg 4560
catactctgc gacatcgtat aacgttactg gtttcacatt caccacctg aattgactct 4620
cttccggggc ctatcatgcc ataccgcgaa aggttttgcg ccattcgatg gtgtccggga 4680
tctcgacgct ctccttatg cgactcctgc attaggaagc agcccagtag taggttgagg 4740


```

ccgttgagca ccgccgccgc aaggaatggt gcatgcaagg agatggcgcc caacagtccc 4800
ccggccacgg ggcttgcac catacccaag ccgaaacaag cgctcatgag cccgaagtgg 4860
cgagcccgat cttcccccac ggtgatgtcg gcgatatagg cgccagcaac cgcacctgtg 4920
gcgccggtga tgccggccac gatgcgtccg gcgtagagga tcgagatctc gatcccgcca 4980
aattaatacg actcactata ggggaattgt gagcggataa caattcccct ctagaaataa 5040
ttttgtttaa ctttaagaag gagatataca tatgcagcat caccaccatc accactggca 5100
gccctcttc ttcaagtggc tcttgctctg ttgccctggg agttctcaaa ttgctgcagc 5160
agcctccacc cagcctgagg atgacatcaa tacacagagg aagaagagtc aggaaaagat 5220
gagagaagtt acagactctc ctgggcgacc ccgagagctt accattcctc agacttcttc 5280
acatggtgct aacagatttg tttgatgaat tctgcagata tccatcacac tggcggccgc 5340
tcgagcacca ccaccaccac cactgagatc cggctgctaa caaagcccga aaggaagctg 5400
agttggctgc tgccaccgct gagcaataac tagcataacc ccttggggcc tctaaacggg 5460
tcttgagggg ttttttgctg aaaggaggaa ctatatccgg at 5502

```

<210> 786

<211> 108

<212> PRT

<213> Homo sapiens

<400> 786

```

Arg Arg Ser Cys Glu Pro Ala Thr Arg Val Pro Glu Val Trp Ile Leu
          5              10              15
Ser Pro Leu Leu Arg His Gly Gly His Thr Gln Thr Gln Asn His Thr
          20              25              30
Ala Ser Pro Arg Ser Pro Val Met Glu Ser Pro Lys Lys Lys Asn Gln
          35              40              45
Gln Leu Lys Val Gly Ile Leu His Leu Gly Ser Arg Gln Lys Lys Ile
          50              55              60
Arg Ile Gln Leu Arg Ser Gln Val Leu Gly Arg Glu Met Arg Asp Met
          65              70              75              80
Glu Gly Asp Leu Gln Glu Leu His Gln Ser Asn Thr Gly Asp Lys Ser
          85              90              95
Gly Phe Gly Phe Arg Arg Gln Gly Glu Asp Asn Thr
          100              105

```

<210> 787

<211> 152

<212> PRT

<213> Homo sapiens

<400> 787

```

Arg Pro Lys Glu Glu Val Pro Arg Ser Lys Ala Leu Glu Val Thr Lys
          5              10              15
Leu Ala Ile Glu Ala Gly Phe Arg His Ile Asp Ser Ala His Leu Tyr
          20              25              30
Asn Asn Glu Glu Gln Val Gly Leu Ala Ile Arg Ser Lys Ile Ala Asp
          35              40              45
Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
          50              55              60
Thr Phe His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Asn Ser Leu
          65              70              75              80
Lys Lys Ala Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Ser Pro

```

```
<210> 788
<211> 1633
<212> DNA
<213> Homo sapiens
```

```
<210> 789
<211> 200
<212> PRT
<213> Homo sapien
```

<400> 789
Met Ala Lys Gly Asp Pro Lys Lys Pro Lys Gly Lys Met Ser Ala Tyr
1 5 10 15
Ala Phe Phe Val Gln Thr Cys Arg Glu Glu His Lys Lys Lys Asn Pro

20 25 30
 Glu Val Pro Val Asn Phe Ala Glu Phe Ser Lys Lys Cys Ser Glu Arg
 35 40 45
 Trp Lys Thr Met Ser Gly Lys Glu Lys Ser Lys Phe Asp Glu Met Ala
 50 55 60
 Lys Ala Asp Lys Val Arg Tyr Asp Arg Glu Met Lys Asp Tyr Gly Pro
 65 70 75 80
 Ala Lys Gly Gly Lys Lys Lys Lys Asp Pro Asn Ala Pro Lys Arg Pro
 85 90 95
 Pro Ser Gly Phe Phe Leu Phe Cys Ser Glu Phe Arg Pro Lys Ile Lys
 100 105 110
 Ser Thr Asn Pro Gly Ile Ser Ile Gly Asp Val Ala Lys Lys Leu Gly
 115 120 125
 Glu Met Trp Asn Asn Leu Asn Asp Ser Glu Lys Gln Pro Tyr Ile Thr
 130 135 140
 Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp Val Ala Asp Tyr
 145 150 155 160
 Lys Ser Lys Gly Lys Phe Asp Gly Ala Lys Gly Pro Ala Lys Val Ala
 165 170 175
 Arg Lys Lys Val Glu Glu Glu Asp Glu Glu Glu Glu Glu Glu Glu
 180 185 190
 Glu Glu Glu Glu Glu Glu Asp Glu
 195 200

<210> 790

<211> 457

<212> DNA

<213> Homo sapiens

<400> 790

ttgcgctgtg ttgggaacgc ggcgagctg tgagccggcg actcgggtcc ctgaggtctg 60
 gattctttct ccgctactga gacacggcgg acacacacaa acacagaacc acacagccag 120
 tcccaggagc ccagtaatgg agagccccaa aaagaagaac cagcagctga aagtcgggat 180
 cctacacctg ggcagcagac agaagaagat caggatacag ctgagatccc agtgcgcgac 240
 atggaaggtg atctgcaaga gctgcatcag tcaaacaccg gggataaatc tggatttggg 300
 ttccggcgtc aaggtgaaga taatacctaa agaggaacac tgtaaaatgc cagaagcagg 360
 tgaagagcaa ccacaagttt aaatgaagac aagctgaaac aacgcaagct ggttttatat 420
 tagatatttg acttaaacta tctcaataaa gttttgc 457

<210> 791

<211> 126

<212> PRT

<213> Homo sapiens

<400> 791

Ser Pro Val Leu Gly Thr Arg Arg Ser Cys Glu Pro Ala Thr Arg Val
 5 10 15

Pro Glu Val Trp Ile Leu Ser Pro Leu Leu Arg His Gly Gly His Thr
 20 25 30

Gln Thr Gln Asn His Thr Ala Ser Pro Arg Ser Pro Val Met Glu Ser
 35 40 45

Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu His Leu Gly
50 55 60

Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln Cys Ala Thr
65 70 75 80

Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro Gly Ile Asn
85 90 95

Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro Lys Glu Glu
100 105 110

His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln Val
115 120 125

<210> 792

<211> 461

<212> DNA

<213> Homo sapiens

<400> 792

cggcggagct gtgagccggc gactcgggtc cctgaggtct ggattctttc tccgctactg 60
agacacggcg gacacacaca aacacagaac cacacagcca gtcccaggag cccagtaatg 120
gagagcccca aaaagaagaa ccagcagctg aaagtcggga tcctacacct ggcgcagcaga 180
cagaagaaga tcaggataca gctgagatcc caggtgctgg gaagggaaat gcgcgacatg 240
gaaggtgatc tgcaagagct gcatcagtca aacaccgggg ataaatctgg atttgggttc 300
cggcgtcaag gtgaagataa tacctaaaaga ggaacactgt aaaatgccag aagcaggtga 360
agagcaacca caagttttaa tgaagacaag ctgaaacaac gcaagctggt tttatattag 420
atatttgact taaactatct caataaagtt ttgcagcttt c 461

<210> 793

<211> 108

<212> PRT

<213> Homo sapiens

<400> 793

Arg Arg Ser Cys Glu Pro Ala Thr Arg Val Pro Glu Val Trp Ile Leu
5 10 15

Ser Pro Leu Leu Arg His Gly Gly His Thr Gln Thr Gln Asn His Thr
20 25 30

Ala Ser Pro Arg Ser Pro Val Met Glu Ser Pro Lys Lys Lys Asn Gln
35 40 45

Gln Leu Lys Val Gly Ile Leu His Leu Gly Ser Arg Gln Lys Lys Ile
50 55 60

Arg Ile Gln Leu Arg Ser Gln Val Leu Gly Arg Glu Met Arg Asp Met
65 70 75 80

Glu Gly Asp Leu Gln Glu Leu His Gln Ser Asn Thr Gly Asp Lys Ser

85

90

95

Gly Phe Gly Phe Arg Arg Gln Gly Glu Asp Asn Thr
 100 105

<210> 794

<211> 970

<212> DNA

<213> Homo sapiens

<400> 794

```

tgggctccca gagctcgggt cctttgcagc ctccaccctg gcgatggctc cctggtccta 60
ctttctctct caaactggct ttttctcatt cctttgactc cgccagactt cctcgcccc 120
atgacctggt gttgtgtctg atcaccccaa cattcctggc tgcccaatgt ggggcaatga 180
agaccccggt gaagggaatgc tagagtgtgt gaaagtggag gacgcacgt caaaggacac 240
ctgaggacgt ctcaaagaag ctcggcggga gagctgagcg ctcggaagaa ccaagaatca 300
tctcttttga aaaatcgatt catcaaata gaatcttctg aacaactgtt caagaaggat 360
gcaaatatca cagtgttaga tgaactttct ggttgacacc tgacaggaag agcctctgta 420
ttggaccacc atgtttgtgc tctgtgtgta gtaacaaacc aacacaccaa aatagcggga 480
gttgccactg acaaagagtt gaatgatcaa atgacggcca aaggaggagg ttccgagaag 540
taaagctttg gaggtcacaa aattagcaat agaagctggg ttccgccata tagattctgc 600
tcatttatac aataatgagg agcagggttg actggccatc cgaagcaaga ttgcagatgg 660
cagtgtgaag agagaagaca tattctacac ttcaaagctt tgggtccactt ttcacgcacc 720
agagttgggt cgaccagcct tggaaaaactc actgaaaaaa gctcaattgg actatgttga 780
cctctatctt attcattctc caatgtctct aaagccagggt gaggaacttt caccaacaga 840
tgaaaatgga aaagtaatat ttgacatagt ggatctctgt accacctggg aggccatgga 900
gaagtgtaag gatgcaggat tggccaagtc cattgggggtg tcaaacttca acccgcaggc 960
agctggagat
  
```

<210> 795

<211> 152

<212> PRT

<213> Homo sapiens

<400> 795

Arg Pro Lys Glu Glu Val Pro Arg Ser Lys Ala Leu Glu Val Thr Lys
 5 10 15

Leu Ala Ile Glu Ala Gly Phe Arg His Ile Asp Ser Ala His Leu Tyr
 20 25 30

Asn Asn Glu Glu Gln Val Gly Leu Ala Ile Arg Ser Lys Ile Ala Asp
 35 40 45

Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
 50 55 60

Thr Phe His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Asn Ser Leu
 65 70 75 80

Lys Lys Ala Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Ser Pro
 85 90 95

Met Ser Leu Lys Pro Gly Glu Glu Leu Ser Pro Thr Asp Glu Asn Gly
 100 105 110

Lys Val Ile Phe Asp Ile Val Asp Leu Cys Thr Thr Trp Glu Ala Met
 115 120 125

Glu Lys Cys Lys Asp Ala Gly Leu Ala Lys Ser Ile Gly Val Ser Asn
 130 135 140

Phe Asn Pro Gln Ala Ala Gly Asp
 145 150

<210> 796

<211> 2435

<212> DNA

<213> Homo sapiens

<400> 796

```
atccactcgg gccgcategc cgcggtgcac aacgtgccgc tgagcgtgct catccggccg 60
ctgccgtccg tgttggaacc cgccaagggt cagagcctcg tggacacgat ccgggaggac 120
ccagacagcg tgccccccat cgatgtcctc tggatcaaag gggcccaggg aggtgactac 180
ttctactcct ttgggggctg ccaccgctac gcggcctacc agcaactgca gcgagagacc 240
atccccgcca agcttgtcca gtccactctc tcagacctaa ggggtgtacct gggagcatcc 300
acaccagact tgcagtagca gcctccttgg cacctgctgc caccttcaag agcccagaag 360
acacacctgg cctccagcag gctgggccat gcagaaggga tagcaggggt gcattctctt 420
tgcacctggc gagaggggtc gactctgggc accctctca ccagctacaa ggccctggac 480
tcaactgtaca gtgtgggagc ccagttccc acctctgtga caataggatc atggccttac 540
ccttgaagca ttaccgagaa ggagaacaga gatgggcttg aagagccacg tgcctgcggc 600
tccaaattcc caaggacaag gatccctctg catttttgtc tatgtaacct cttatatgga 660
ctacattcag ctgcaaggaa aggaaaacct tgattgcagt ggtttaaaca aacagaagat 720
tgtttttcca catagcatgg attctggaga tgggtggcta atggtattgg ttcaacaact 780
ccacgaaggt aggggtcacg tcttggatcc ttttgctta atctcagtgc tcgttacttc 840
atggtcccaa gatggctgct gtatcccaa gaatcatgtc tgcgttcaag gaaggagggg 900
tgagggaaga ggaaggcca aactagctgg acccgtcacc ttctatcaga aagtaaaacc 960
tcgtcagaag tctgtttcct gctctctccc tctgcatac ttcaactaga tgcccttggc 1020
ccgagccagc taccattgca cctctagctg caaacaagc taagacagca gggaacagaa 1080
ttgtcatggc tgaatagacc aatcgtgttc catctactga gactggcaca ctgcctcctg 1140
caataaaact gggatcccat taccaagaga gaaatgcaga attgtgtacc agttagcttt 1200
tgctgtgtaa caaaccatcc ccaaacttgg cagctagaaa caaacctgt attttccac 1260
aatcctatgg gttggcaatt tgggctgggc tcaacagggc agttctgctg ctcacacctg 1320
ggatccctca tggagctaag gtcagctgtt acctcagctg ggctggatg gtctaggata 1380
gccttactca cttgcctggc aggtgacagg ctgttggctg gaattgcttg gttctcctcc 1440
atgtggcctc tccagcaggc tagctcaggc ttattcacat gatggcttca ggattccaaa 1500
gagagtgaga gtagaagctg aaagacttct tgagttcttg gcctggaact gggactagga 1560
cagtgtcact tctgctaagt tcttttggtc agagcaaata acaaggcttt acccagattc 1620
aagggatgag aaacagacta catgtcttga tgaggggaac cacaaaagc ttgtggccat 1680
ttttcaccta tcacaaataa ttttggatgg gtatttattt ggataaaggt atttccctct 1740
tcccccttct cctctgtctc atggggcctc actctgcca gttggaaggc actaagacat 1800
tgtcctggcc tctagggctc aggggaagag gtgttggggc aggaagtga tctctccatg 1860
ggctggaccc actgtagttag gagtgccctc ttgtctgcac tgctggtagt gggttaggcc 1920
aggtaggaca ttccagaggg gcttctgaaa accaagagtc cctggggaaa gggaacagag 1980
taaggcaggc cttgtttctc ctgccctcta agggaacttg gtcactcggc acttttaagc 2040
```



```
<210> 799
<211> 60
<212> PRT
<213> Homo sapiens
```

```
<210> 800
<211> 2477
<212> DNA
<213> Homo sapien
```

```
<400> 800
gccttgccaa aaaagcacia gccaaccca gccctgattg ccctgcgcta ccagctacag 60
cgtgggggttg tggtcctggc caagagctac aatgagcagc gcatcagaca gaacgtgcag 120
```


gtgtttgaat tccagttgac ttcagaggag atgaaagcca tagatggcct aaacagaaat 180
 gtgcgatatt tgacccttga tatttttgcg gggcccccta attatccatt ttctgatgaa 240
 tattaacatg gagggcattg catgaggtct gccagaaggc cctgcgtgtg gatggtgaca 300
 cagaggatgg ctctatgctg gtgactggac acatcgctc tggttaaatc tctcctgctt 360
 ggtgatttca gcaagctaca gcaaagccca ttggccagaa aggaaagaca ataattttgt 420
 tttttcattt tgaaaaaatt aaatgctctc tcctaaagat tcttcaccta ctttggctctc 480
 cataacttct atgttttctt tccttctgac acactagtgc ccctaaattg tgatttgcct 540
 atacgtttag ggccgggggtt ggaagatggt aacaaccatt taagattcat ttctgcagt 600
 ggagtgggtg gagtttcacc ctctgggaaa ggggcagggtg acaggatatt atcagtcagt 660
 gcctctctag ctcttgtagg aagaagcaca cgcaggatgg agtctagagg atgagcgata 720
 ttgactagca attcatgggc tccctccagc agtgcgaggg tcagagtttc tggagccttg 780
 ggaggaggca tccctgtgag ggggggttag ggagatggga gggcaccagg aaaagtgtatt 840
 agaagtcagg tatgggaagg cttaaataagga cagagtcgag tacatctctg cttggaaaaa 900
 catatcaaca cccttttttt tgaacattat atcttgctca taaaagaaaa ctttccacat 960
 tgttttaaca aacccacag ctgagagtca ggcctgaatc tttgatgtgt gccagtcac 1020
 agagttgacc ctattgggtt gtggtggggc agggcatcaa agacatcatt gactaatcac 1080
 attcccctga atagctcata tttagaaaaat attcttagat tctaaaaatg tactattaat 1140
 ttgtgatatt cagtctttta aatattttat acattaaaca ggcatagtta caaatataaa 1200
 acaaaaaat cccaaagcca ttatgcatgg cactcaagat taaaatggga aataatacat 1260
 ctaataaatc aaatgttcca agacttcaaa ggtcttttgg aaacaggcta tgtaaaacag 1320
 cacactggtt tcaaactttg gtaaatttta agaacaactc ttacaaaggc atttaattct 1380
 tatacataat tttcagggga cctaagttaa tcagctaate atgaagacat gattttcatt 1440
 ttagaaaaa cttttgaaaa cttgggataa tctcatgcct taatgatcaa agcattatga 1500
 gaaggacagt ggtttttaac ctgggcatat gttctaacac atttactctc cactattcgt 1560
 actctggtag ccagtgttaac cccatcagag attccttctc aagccatgtc tcagagctga 1620
 gaggcacccc agcaagtttt gcagctcaca gttttttccg taaattactt attctataaa 1680
 attggagtag gccataaact ttggagggcc ctagaccaat tttttggatt atttttcgtc 1740
 ttctatcatt ccgctgatct tagatattct ctgcattaaa tattaatat cacttctagg 1800
 ctgaaaaatc ccctaaaaa tatttctagc tcagattttt cctccaaatt ctgcaataga 1860
 agatcacaat gtgaactctg catctccatg ttaaagtcta atggacattc acacttagca 1920
 tgtctcaaag aaatctcatg taaaccatgg ccactctgtt ctaccttaac tttctgagtc 1980
 tatggaatga taatttcaca tctcataaac ttgactgatg taagtgtcaa gaaaagattg 2040
 acattttgtt aaaagttagt agtgaagtgt gtaacgctta agcaaaactt catatttcaa 2100
 atctctttag caagtgtaac tcttttttca agatgtgaaa taatcattag gtcagtcatt 2160
 tgtaaatagt acatctgcta tggacttttt ccagttcttc accatccatt tttataaaac 2220
 tcttattgtt aaaaaaaaag ttactcagaa tttcataaag ccaaacacct gatttcagga 2280
 acacttgaga tgtaagaaaa ttttataggg acctccaatc actaattttc ctattttttc 2340
 tctcaaagaa atgctgaagg gaggaattca ggttgaatga aaggaaatag taacttacag 2400
 ccatatagag ttataaagac ttcttgtaaa tgtgaacata tggtaaaata taaaaacatg 2460
 tatttttgaa aaaaaaa 2477

<210> 801
 <211> 1619
 <212> DNA
 <213> Homo sapien

<400> 801
 ggtacgcgcc cgcttgcgct ccggcctcta ctgcggcggtc atcgtctacg acgagcgcag 60
 cccgcgcgcc gagagcctcc gcgaggacag caccgtgtcg ctggtggtgc aggcgctgcg 120
 ccgcaacgcc gagcgcaccg acatctgcct gctcaaaggc ggctatgaga ggttttccctc 180
 cgagtaccca gaattctgtt ctaaaaccaa ggccttggca gccatccac ccccggttcc 240
 cccagtgcc acagagccct tggacctggg ctgcagctcc tgtgggaccc cactacacga 300
 ccaggggggt cctgtggaga tccttccctt cctctacctc ggagtgctc accatgctgc 360
 ccggagagac atgctggacg ccctgggcat cacggctctg ttgaatgtct cctcggaactg 420

```

cccaaaccac tttgaaggac actatcagta caagtgcac ccagtgggaag ataaccacaa 480
ggccgacatc agctcctggt tcatggaagc catagagtac atcgatgccg tgaaggactg 540
ccgtgggcgc gtgctggtgc actgccaggc gggcatctcg cggtcggcca ccatctgcct 600
ggcctacctg atgatgaaga aacgggtgag gctggaggag gccttcgagt tcgttaagca 660
gcgcccgcgc attatctcgc ccaacttcag cttcatgggg cagctgctgc agttcgagtc 720
ccaggtgctg gccacgtcct gtgctgcgga ggctgctagc ccctcgggac ccctgcggga 780
gcggggcaag acccccgcga cccccacctc gcagttcgtc ttcagctttc cggctctccgt 840
gggctgacac tcggcccccga gcagcctgcc ctacctgcac agcccatca ccacctctcc 900
cagctgttag agccgccctg ggggccccag aaccagagct ggctcccagc aagggttagga 960
cgggcgcgat gcgggcagaa agttgggact gagcagctgg gagcaggcga ccgagctcct 1020
tccccatcat ttctccttgg ccaacgacga ggccagccag aatggcaata aggactccga 1080
atacataata aaagcaaaaca gaacactcca acttagagca ataacggctg ccgcagcagc 1140
cagggaagac cttggtttgg tttatgtgtc agtttcactt ttccgataga aatttcttac 1200
ctcatttttt taagcagtaa ggcttgaagt gatgaaaccc acagatccta gcaaatgtgc 1260
ccaaccagct ttactaaagg gggaggaagg gagggcaaa ggatgagaag acaagtttcc 1320
cagaagtgcc tggttctgtg tacttgtccc tttgtgtcgt ttgtttagt taaaggaatt 1380
tcatttttta aaagaaatct tcgaagggtt ggttttcatt tctcagtcac caacagatga 1440
ataattatgc ttaataataa agtatttatt aagactttct tcagagtatg aaagtacaaa 1500
aagtctagtt acagtggatt tagaatatat ttatgttgat gtcaaacagc tgagcaccgt 1560
agcatgcaga tgtcaaggca gttaggaaga attaggtttg aattgctttt taaaaaaaa 1619

```

<210> 802

<211> 3115

<212> DNA

<213> Homo sapien

<400> 802

```

cgctccgcga cgcgtgggct catcttgaga agcaggcggg ttgggtggga ggaggaagaa 60
agggagaaga taggtttgaa ttgctttttt aaaaaaaaaa aaaagaaaaa aaaagacagc 120
atctcactat gttgccaaagg ctcatctcaa gctcttgggc tcaagagatc ctcccacctc 180
ggcctcctga gtagctggga ctgcagggtg gtgtcatcat gaccaatgtg aattgctttt 240
gaagctgggt catgggcctg taggccaccg aagcaatttt agaccacagt aagtcaagct 300
tttttccctc cgatgatcac tgggtggttg cagcattttt tgcataaaacc tgcctaagac 360
ttgtctatcg tctgtgatca atatgccata ttacactaag gtgctcctgg aaaattgggt 420
gcagttcaaa ttttctaca gcaaatcatt tggcaaggcc agccattggg gaaaccagac 480
aactagagat aaccctgaaa tgaatccttt tgtaaattga agcaccatct tttctttttt 540
tgcataaatt ggaggtttta attttagggc agttacctga agtgaaatat accaacaatt 600
tcttgtgttc tttaaattcc tagttagggt aatatttttg aaggtcctct tttgaataaa 660
gaggggaatg gacaccacat ttcaggtcct ctggaagtgt ggaagggcaa gagagcatca 720
gtgagctgat ggtggattgc ttacatcgga ttccattggg atgaatttcc caaactggaa 780
atcaaagcgc cagggtgggg ttggggctga ctgctggtga gggggctggc cgctggctcc 840
cgtgacgtgc gtcatgggca cgcaggcgcc attttgaatc tatcgtcggc acgtgggtgc 900
cattttgaat ccttagttgg gcctttctaa atggagaatg gctttggagg gagacacgtt 960
ttctgtgggg agggtttggg ggggagggag gagggaaaca gctacatgct attttgtttg 1020
tagtattgtg gaacagtctt gttatggagt gccagcttag aggttgttgc aaacttgtct 1080
agaagtgaga gcatggtttt ttttagccct ttgagagtct acatctaata aacattcttg 1140
ctcaccataa aataacgtca agcctcaatg tcaccgtcac gttgggatac tctttctcat 1200
ctggcatcct agacaggaca aggttggtta cctttccttc catgaaccat gaacctgtga 1260
cggcatcatt catcctgact tcaccaagct cgcctgtgg gtgaggccag agtcccact 1320
ggcaattttt agaagagcca gaggtccctt gcttcctcta gaaataacag ttcagggtga 1380
agcatggagg gtttcagttc ccagacaatg gaaccattta gagacaacac agttggacat 1440
ttccactttt tccttgatcc ctggaagtcc agtgggttct gcagctgaaa aagccctggg 1500
tcccagcagc agagagacag gacagagggg atgcttgggc ggggaggggc ggtaacctgc 1560

```

agaacagatt ccatttttat agaacgagta cacgtttgct aaaacagtcc tgctttccca 1620
 gactggattc ccaccacagg gacagtccga actcaggact agctccagcg acatctttcc 1680
 tccgaattca agccttctat cacaatgtca aaacagctat ttataaagcc attttcattg 1740
 tacttgataa cagcacgagt cccaaaactt ttagaaataa aataggacat tggcttgatt 1800
 gaaaagaggg acttttttaa aattgttctt tcgtcagaag ccttttggat gacttacaat 1860
 agctctgatg aagataccac ccagcgtca gtccaatagg tcagtgaagt tcaacaggca 1920
 tccatccctc ccatgaaggg attctggtga ggggaagttt ctgtaatgac aggaaagcat 1980
 tgaccctcat tgattgtcaa ctttgggtatt agccatgaaa gacaggatgc tcattgggtg 2040
 ttctgtagag tgaggaatgc tgccatttcc ctcccagAAC gtctgaccca ggggtgtgtg 2100
 ttgaggagcc ctgggggaaa tggaccaagt tttcccacag agcagtatta ggctgaagag 2160
 caggtgactg gtaggccccca gctcccatca tttccctccca aagccatttt gttcagttgc 2220
 tcatccacgc tggattccag agagttttcc aatttgggaa gccatgagaa aggtttttta 2280
 atcttgggaa gatggagaga gggacatagg atagttgact ccaacatgac aggaagaggc 2340
 tggagattgg gaattggcca tcaaccaagc ctgtagtagt aaagccatgg tcccgcattg 2400
 gaattacttg gggaacttat acagtctctga taccagggtc ctcttagacc agttcaacca 2460
 attctaggtg ggggactcag gcatcagtggt gtttcgttagc tcccgggtg ttttccctgt 2520
 gcagccgagc ttgggaaact gccatgcttt ttggatgtca aggcgctgtt ggaggctggg 2580
 tgtgacagca cagagccagg ttgtcttggt gaaaccacag ccacgggttt gccactggct 2640
 cagcatggcc tcactgccag tcccagcctg gctgagggac aagatgggtt ctcttgggag 2700
 ttctctagtg gagcaccctt ccaggctttt tgaaagccag ctgatctgtg gagccttgtt 2760
 aagggactca atacggtgtt tggatattga tgtttttcct tgagactgtc ttgtccatca 2820
 ataaagatgg aggatgtctc ctctttgaac cccgcttccc caccagtact ctctctccct 2880
 tagagtttat gagttattca aggaggagac ttcttaaaga cagcaacgca attcttgtaa 2940
 cttgtgtaaa tagcccatc tttcagagtg ataccatttc tacatttgat aatgcctgta 3000
 ttctgttagg atgtatatag tttaggggat tttttttttg tttgggtttg ttttttagaa 3060
 gtcaatatgt ctggttttat ttattgcttg aaaaagatca tttgaaaaaa ataaa 3115

<210> 803
 <211> 1238
 <212> DNA
 <213> Homo sapien

<400> 803
 cccgggttct cttctcttcc tcgcgcgcc agccgcctcg gttcccggcg accatggtga 60
 cgatggagga gctgcgggag atggactgca gtgtgctcaa aaggctgatg aaccgggacg 120
 agaattggcg cggcgcgggc ggcagcggca gccacggcac cctggggctg ccgagcggcg 180
 gcaagtgcct gctgctggac tgcagaccgt tcttggcgca cagcgcgggc tacatcctag 240
 gttcgggtcaa cgtgcgctgt aacaccatcg tgcgcggcg ggctaagggc tccgtgagcc 300
 tggagcagat cctgcccgcc gaggaggagg tacgcgcccg cttgcgctcc ggccctctact 360
 cggcgggtcat cgtctacgac gagcgcagcc cgcgcgccga gagcctccgc gaggacagca 420
 ccgtgtcgct ggtggtgcag gcgctgcgcc gcaacgccga gcgcaccgac atctgcctgc 480
 tcaaaggcgg ctatgagagg ttttccctcc agtaccacga attctgttct aaaaccaagg 540
 ccttggcagc catcccaccc ccggttcccc ccagcggcac agagcccttg gacctggact 600
 gcagctcctg tgggacccca ctacacgacc aggagggtcc tgtggagatc cttcccttcc 660
 totacctcgg cagtgcctac catgctgccc ggagagacat gctggacgcc ctgggcatca 720
 cggtctgtgt gaattgtctc tcggactgcc caaaccactt tgaaggacac tatcagtaca 780
 agtgcacccc agtggaagat aaccacaagg ccgacatcag ctcttggttc atggaagcca 840
 tagagtacat cgatgccgtg aaggactgcc gtgggcgct gctggtgcac tgccaggcgg 900
 gcatctcgcg gtcggccacc atctgcctgg cctacctgat gatgaagaaa cgggtgaggc 960
 tggaggaggc cttcagattc gtttaagcag gccgcagcat catctcgccc aacttcagct 1020
 tcatggggca gctcgtgcag ttcgagtccc aggtgctggc cacgtcctgt gctgcggagg 1080
 ctgctagccc ctcgggaccc ctgggggagc ggggcaagac ccccgccacc cccacctcgc 1140
 agttcgtctt cagctttccg gtctccgtgg gcgtgcactc ggccccagc agcctgccct 1200
 acctgcacag ccccatcacc acctctccca gctgttag 1238

<210> 804
 <211> 4637
 <212> DNA
 <213> Homo sapiens

<400> 804

```

ggtacgcgcc cgcttgccgt cccgcctcta ctccggcggtc atcgtctacg acgagcgcag 60
cccgcgcgcc gagagcctcc gcgaggacag caccgtgtcg ctgggtgggtc aggcgctgcg 120
ccgcaacgcc gagcgcaccg acatctgcct gctcaaaggc ggctatgaga ggttttcctc 180
cgagtaccga gaattctgtt ctaaaaccaa ggccctggca gccatcccac ccccggttcc 240
ccccagtgcc acagagccct tggacctggg ctgcagctcc tgtgggacct cactacacga 300
ccaggggggt cctgtggaga tccttccctt cctctacctc ggcagtgcct accatgctgc 360
ccggagagac atgctggacg ccctgggcat caccgctctg ttgaatgtct cctcggactg 420
cccaaaccac tttgaaggac actatcagta caagtgcac ccagtgaag ataaccacaa 480
ggccgacatc agctcctggt tcatggaagc catagagtac atcgatgccg tgaaggactg 540
ccgtggggcg gtgctgggtc actgccaggc gggcatctcg cggtcggcca ccatctgcct 600
ggcctacctg atgatgaaga aacgggtgag gctggaggag gccttcgagt tcgttaagca 660
gcgcgcgacg attatctcgc ccaacttcag cttcatgggg cagctgctgc agttcgagtc 720
ccaggtgctg gccacgtcct gtgctgcgga ggctgctagc ccctcgggac ccctgcggga 780
gcggggcaag acccccgcga cccccacctc gcagttctgc ttcagctttc cggctctcgt 840
gggctgacac tcggcccccga gcagcctgcc ctacctgcac agcccccata ccacctctcc 900
cagctgttag agccgccttg ggggcccag aaccagagct ggctcccagc aagggttagga 960
cgggcccgat gcgggcagaa agttgggact gagcagctgg gagcaggcga ccgagctcct 1020
tccccatcat ttctccttgg ccaacgacga ggccagccag aatggcaata aggactccga 1080
atacataata aaagcaaaac gaacactcca acttagagca ataacggctg ccgcagcagc 1140
cagggaagac cttggttttg tttatgtgtc agtttcaact ttccgataga aatttcttac 1200
ctcatttttt taagcagtaa ggcttgaagt gatgaaaacc acagatccta gcaaatgtgc 1260
ccaaccagct ttactaaagg gggaggaagg gagggcaaa ggaatgagaag acaagtttcc 1320
cagaagtgcc tggttctgtg tacttgtccc tttgttgcg ttgtttagt taaaggaatt 1380
tcatttttta aaagaaatct tcgaaggtgt ggttttcatt tctcagtcac caacagatga 1440
ataattatgc ttaataataa agtatttatt aagactttct tcagagtatg aaagtacaaa 1500
aagtctagtt acagtggatt tagaatatat ttatgttgat gtcaaacagc tgagcaccgt 1560
agcatgcaga tgtcaaggca gttaggaaga attaggtttg aattgctttt ttaaaaaaaa 1620
agaaaagaaa aaaaaagaca gcattctact atgttgccaa ggctcatctc aagctcttgg 1680
gctcaagaga tcctccacc tcggcctcct gagtagctgg gactgcaggt gtgtgtcatc 1740
atgaccaatg tgaattgctt ttgaagctgg ttcatgggca ttagggccac cgaagcaatt 1800
ttagaccaca gtaagtcaag cttttttccc tccgatgatc actgggtggg tgcagcattt 1860
tttgcataaa cctgcctaag acttgtctat cgtctgtgat caatatgcca tattacacta 1920
aggtgctcct ggaaaatttg gtgcagttca aattttccta cagcaaatca tttggcaagg 1980
ccagccattg gggaaaccag acaactagag ataaccctga aatgaatcct tttgtaaatt 2040
gaagcaccat cttttctttt tttgcataaa ttggaggttt taatttttagg gcagttacct 2100
gaagtgaat ataccaacaa tttcttgtgt tctttaaatt cctagttagg tgaatatatt 2160
tgaaggctct cttttgaata aagaggggaa tggacaccac atttcagggt ttctcgaagt 2220
gtggaagggt aagagagcat cagtgcagct atgggtggat gcttacatcg gattccattg 2280
gtatgaattt cccaaactgg aaatcaaagc gccagggtgg ggttggggct gactgctggt 2340
gagggggctg gccgctgggt cccgtgacgt gcgtcatggg cacgcaggcg ccattttgaa 2400
tctatcgtcg gcacgtgggt gccattttga atccttagtt gggcctttct aaatggagaa 2460
tggctttgga gggagacacg ttttctgttg ggaagggttg ggggggaggg aggaggggaa 2520
aagctacatg ctattttgtt ttagtatttg ttgaacagtc ttgttatgga gtgccagctt 2580
agaggttgtt gcaaacttgt ctagaagtga gagcatggtt ttttttagcc ctttgagagt 2640
ctacatctaa tgaacattct tgctcaccga taaataacgt caagcctcaa tgtcaccgtc 2700
acgttgggat actctttctc atctggcatc ctgacagga caagggtggg tacctttcct 2760
tccatgaacc atgaacctgt gacggcatca ttcactctga cttaccaag ctccgcctgt 2820

```

```
210> 805
<211> 394
<212> PRT
<213> Homo sapiens
```

Arg Leu Met Asn Arg Asp Glu Asn Gly Gly Gly Ala Gly Gly Ser Gly
20 25 30

Ser His Gly Thr Leu Gly Leu Pro Ser Gly Gly Lys Cys Leu Leu Leu
35 40 45

Asp Cys Arg Pro Phe Leu Ala His Ser Ala Gly Tyr Ile Leu Gly Ser
50 55 60

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Asn | Val | Arg | Cys | Asn | Thr | Ile | Val | Arg | Arg | Arg | Ala | Lys | Gly | Ser |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |

Val Ser Leu Glu Gln Ile Leu Pro Ala Glu Glu Glu Val Arg Ala Arg

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|--|
| | | | | 85 | | | | | 90 | | | | | 95 | | | |
| Leu | Arg | Ser | Gly | Leu | Tyr | Ser | Ala | Val | Ile | Val | Tyr | Asp | Glu | Arg | Ser | | |
| | | | 100 | | | | | 105 | | | | | 110 | | | | |
| Pro | Arg | Ala | Glu | Ser | Leu | Arg | Glu | Asp | Ser | Thr | Val | Ser | Leu | Val | Val | | |
| | | 115 | | | | | 120 | | | | | 125 | | | | | |
| Gln | Ala | Leu | Arg | Arg | Asn | Ala | Glu | Arg | Thr | Asp | Ile | Cys | Leu | Leu | Lys | | |
| | 130 | | | | | 135 | | | | | 140 | | | | | | |
| Gly | Gly | Tyr | Glu | Arg | Phe | Ser | Ser | Glu | Tyr | Pro | Glu | Phe | Cys | Ser | Lys | | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | | |
| Thr | Lys | Ala | Leu | Ala | Ala | Ile | Pro | Pro | Pro | Val | Pro | Pro | Ser | Ala | Thr | | |
| | | | | 165 | | | | | 170 | | | | | 175 | | | |
| Glu | Pro | Leu | Asp | Leu | Asp | Cys | Ser | Ser | Cys | Gly | Thr | Pro | Leu | His | Asp | | |
| | | | 180 | | | | | 185 | | | | | 190 | | | | |
| Gln | Glu | Gly | Pro | Val | Glu | Ile | Leu | Pro | Phe | Leu | Tyr | Leu | Gly | Ser | Ala | | |
| | | 195 | | | | | 200 | | | | | 205 | | | | | |
| Tyr | His | Ala | Ala | Arg | Arg | Asp | Met | Leu | Asp | Ala | Leu | Gly | Ile | Thr | Ala | | |
| | 210 | | | | | 215 | | | | | 220 | | | | | | |
| Leu | Leu | Asn | Val | Ser | Ser | Asp | Cys | Pro | Asn | His | Phe | Glu | Gly | His | Tyr | | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | | |
| Gln | Tyr | Lys | Cys | Ile | Pro | Val | Glu | Asp | Asn | His | Lys | Ala | Asp | Ile | Ser | | |
| | | | | 245 | | | | | 250 | | | | | 255 | | | |
| Ser | Trp | Phe | Met | Glu | Ala | Ile | Glu | Tyr | Ile | Asp | Ala | Val | Lys | Asp | Cys | | |
| | | | 260 | | | | | 265 | | | | | 270 | | | | |
| Arg | Gly | Arg | Val | Leu | Val | His | Cys | Gln | Ala | Gly | Ile | Ser | Arg | Ser | Ala | | |
| | | 275 | | | | | 280 | | | | | 285 | | | | | |
| Thr | Ile | Cys | Leu | Ala | Tyr | Leu | Met | Met | Lys | Lys | Arg | Val | Arg | Leu | Glu | | |
| | 290 | | | | | 295 | | | | | 300 | | | | | | |
| Glu | Ala | Phe | Glu | Phe | Val | Lys | Gln | Arg | Arg | Ser | Ile | Ile | Ser | Pro | Asn | | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | | |
| Phe | Ser | Phe | Met | Gly | Gln | Leu | Leu | Gln | Phe | Glu | Ser | Gln | Val | Leu | Ala | | |
| | | | | 325 | | | | | 330 | | | | | 335 | | | |
| Thr | Ser | Cys | Ala | Ala | Glu | Ala | Ala | Ser | Pro | Ser | Gly | Pro | Leu | Gly | Glu | | |
| | | | 340 | | | | | 345 | | | | | 350 | | | | |
| Arg | Gly | Lys | Thr | Pro | Ala | Thr | Pro | Thr | Ser | Gln | Phe | Val | Phe | Ser | Phe | | |
| | | 355 | | | | | 360 | | | | | 365 | | | | | |
| Pro | Val | Ser | Val | Gly | Val | His | Ser | Ala | Pro | Ser | Ser | Leu | Pro | Tyr | Leu | | |

370 375 380

His Ser Pro Ile Thr Thr Ser Pro Ser Cys
385 390

<210> 806
<211> 302
<212> PRT
<213> Homo sapiens

<400> 806
Val Arg Ala Arg Leu Arg Ser Gly Leu Tyr Ser Ala Val Ile Val Tyr
5 10 15
Asp Glu Arg Ser Pro Arg Ala Glu Ser Leu Arg Glu Asp Ser Thr Val
20 25 30
Ser Leu Val Val Gln Ala Leu Arg Arg Asn Ala Glu Arg Thr Asp Ile
35 40 45
Cys Leu Leu Lys Gly Gly Tyr Glu Arg Phe Ser Ser Glu Tyr Pro Glu
50 55 60
Phe Cys Ser Lys Thr Lys Ala Leu Ala Ala Ile Pro Pro Pro Val Pro
65 70 75 80
Pro Ser Ala Thr Glu Pro Leu Asp Leu Gly Cys Ser Ser Cys Gly Thr
85 90 95
Pro Leu His Asp Gln Gly Gly Pro Val Glu Ile Leu Pro Phe Leu Tyr
100 105 110
Leu Gly Ser Ala Tyr His Ala Ala Arg Arg Asp Met Leu Asp Ala Leu
115 120 125
Gly Ile Thr Ala Leu Leu Asn Val Ser Ser Asp Cys Pro Asn His Phe
130 135 140
Glu Gly His Tyr Gln Tyr Lys Cys Ile Pro Val Glu Asp Asn His Lys
145 150 155 160
Ala Asp Ile Ser Ser Trp Phe Met Glu Ala Ile Glu Tyr Ile Asp Ala
165 170 175
Val Lys Asp Cys Arg Gly Arg Val Leu Val His Cys Gln Ala Gly Ile
180 185 190
Ser Arg Ser Ala Thr Ile Cys Leu Ala Tyr Leu Met Met Lys Lys Arg
195 200 205
Val Arg Leu Glu Glu Ala Phe Glu Phe Val Lys Gln Arg Arg Ser Ile
210 215 220

Ile Ser Pro Asn Phe Ser Phe Met Gly Gln Leu Leu Gln Phe Glu Ser
 225 230 235 240

Gln Val Leu Ala Thr Ser Cys Ala Ala Glu Ala Ala Ser Pro Ser Gly
 245 250 255

Pro Leu Arg Glu Arg Gly Lys Thr Pro Ala Thr Pro Thr Ser Gln Phe
 260 265 270

Val Phe Ser Phe Pro Val Ser Val Gly Val His Ser Ala Pro Ser Ser
 275 280 285

Leu Pro Tyr Leu His Ser Pro Ile Thr Thr Ser Pro Ser Cys
 290 295 300

<210> 807
 <211> 3829
 <212> DNA
 <213> Homo sapiens

<400> 807
 gtttgaaagt gtgtagcacc tccaccttct ctctctctct cctctccct ctctgccag 60
 ccaagtgaag acatgcttac ttccccttca ccttccttca tgatgtggga agagtgcctgc 120
 aacccagccc tagccaacgc cgcattgagag ggagtgtgcc gagggcttct gagaagggtt 180
 ctctcacatc tagaaagaag cgcttaagat gtggcagccc ctctcttcca agtggctctt 240
 gtctgtttgc cctgggagtt ctcaaattgc tgcagcagcc tccacccagc ctgaggatga 300
 catcaatata cagaggaaga agagtcagga aaagatgaga gaagttacag actctcctgg 360
 gcgaccccgga gagcttacca ttctcagac ttcttcacat ggtgctaaca gatttgcttc 420
 taaaagtaaa gctctagagg ccgtcaaatt ggcaatagaa gccgggttcc accatattga 480
 ttctgcacat gtttacaata atgaggagca ggttgactg gccatccgaa gcaagattgc 540
 agatggcagt gtgaagagag aagacatatt ctacacttca aagctttgga gcaattccca 600
 tcgaccagag ttggtccgac cagccttgga aaggtcactg aaaaatcttc aattggacta 660
 tgttgacctc tatcttattc attttccagt gtctgtaaag ccaggtgagg aagtgatccc 720
 aaaagatgaa aatggaaaaa tactatttga cacagtggat ctctgtgcca catgggaggc 780
 catggagaag tgtaaagatg caggattggc caagtccatc ggggtgtcca acttcaacca 840
 caggctgctg gagatgatcc tcaacaagcc agggctcaag tacaagcctg tctgcaacca 900
 ggtggaatgt catccttact tcaaccagag aaaactgctg gatttctgca agtcaaaaaga 960
 cattgttctg gttgctata gtgctctggg atcccatcga gaagaaccat ggtgggaccc 1020
 gaactccccg gtgctcttgg aggacccagt cctttgtgcc ttggcaaaaa agcacaagcg 1080
 aacccagccc ctgattgccc tgcgctacca gctgcagcgt ggggttgtgg tcctggccaa 1140
 gagctacaat gagcagcgca tcagacagaa cgtgcagggtg tttgaattcc agttgacttc 1200
 agaggagatg aaagccatag atggcctaaa cagaaatgtg cgatatttga ccttgatat 1260
 ttttgctggc ccccttaatt atccattttc tgatgaatat taacatggag ggcattgcat 1320
 gaggtctgcc agaagccct gcgtgtggat ggtgacacag aggatggctc tatgctggtg 1380
 actggacaca tcgctcttgg ttaaactctc cctgcttggc gacttcagta agctacagct 1440
 aagcccatcg gccggaaaag aaagacaata attttgtttt tcattttgaa aaaattaaat 1500
 gctctctcct aaagattctt cactacttt gtctccata acttctatgt tttctctcct 1560
 tctgacacac tagtgcccc aaattgtgat ttgcctatac gtttagggcc gggattggaa 1620
 gatgtaaca accatttaag attcatttct gcagtgggag tgggtggagt ttcacctct 1680
 gggaaagggg caggtgacag gtatttatca gtcagtgcct ctctagctct tgtaggaaga 1740
 agcacacgca ggatggagtc tagaggatga gcgatattga ccagcaattc atgggctccc 1800
 tccagcagtg cgagggtcag agtttctgga gccttgggag gaggcaacce tgtgaggggg 1860

ggtaggggag atgggagggc accaggaaaa gtgattagaa gtcaggtatg ggaaggctaa 1920
 ataggacaga gtcaggtaca tctctgcttg gaaaaacata tcaacaccct ttttttttga 1980
 tcattatatac ttgttcataa aagaaaactt tccacattgt ttttaacaaac cccacagctg 2040
 agagtcaggc ctgaatcttt gatgtgtgcc cattcacaac gttgacccta ttggtttgtg 2100
 gtggggcagg acatcgaaga tatcattgac taatcacatt cccctgaata gctcatatct 2160
 agaaaatatt cttagattgt aaaaatgtac tgttcatttg ttatattcaa tcttttaaat 2220
 gttttatact ttaaacaagg catagttaca agtataaaac ataaatatcc caaagccatt 2280
 atgcatggca ctcaagatta aaatgggaaa taatacatct aataaatcaa atgttccaag 2340
 acttcaaatg tcttttgga acaggctatg taaaacagca cactggtttc aaactttggg 2400
 aaattttaag aagaactctt acaaaggcat ttaattctta tacataattt tcaggggacc 2460
 taagttaatc agctaatac gaagacatga ttttcgtttt agaaaacact tttgaaaact 2520
 tgggataatc tcatgtctta atgatcaaag cattatgaga aggacagtgg ttttttacct 2580
 gggcacactt tctaacacat ttactctcca ctattcgtac tctggtagcc acgttaaccc 2640
 catcagagat tccttctcaa gccatgtctc agagctgata ggcattcccag caagttttgc 2700
 agtcacaat ttttctgtaa attacttatt ctataaaatt ggaagaggcc ataaactttg 2760
 gagggcccta gaccaatttt ttggattatt tctgggtctac tctcattccg ttgatgatct 2820
 tagatattct ctgcattaaa tatcacctct aggctgagaa atccacaaa aaatatttct 2880
 agctcagcgt tttcctccaa atcttcaatg gaagatcata atgtgaactc tgcattctca 2940
 tgttaaagtt taatggacat tcacatttag catgtctcaa agaatctca tgtaaaccat 3000
 ggccatcctg ttctacctta actttctgag tctatggaat gataatttca catctcataa 3060
 acttgactga tgtaagtgtc aagaaaagat tgacattttg ttaaaacttc gtagccaagt 3120
 gtgtaacgct taagcagact ttcatatttc aaatctctat agcacgtgta actctttttt 3180
 caagatgtga aataatcatt aggtcagtc tttgtaaata gtacagctgc tgtgggcttt 3240
 ttccagttct tcaccatcca tttttataaa actcttattg ttaaaaaaaa aaagttactc 3300
 agaatttcat aaagccaaac acctgatttc aggaacactt gagatgtaag aaaattttat 3360
 agggacctcc aatcactaat tttctatttt tttctctcaa agaatgctg aaggagggaa 3420
 ttcaggttga atgaaaggaa atagtaactt acagccatat agagttataa agacttcttg 3480
 taaatgtgaa catatggtaa aatataaaaa catgtatttt tgaaaaaatg gattctactc 3540
 attattttac ttccatttaa gatataaatg tagagaaata agtataattc taagctaata 3600
 cgtacgcaat gtaggaagct gtaattactg accaaaacta tgtgaagtgg agaaaacctg 3660
 gggaagtgga tggttttaga tgaaactgaa gttaaaattc tattgattta aagtaattg 3720
 ttataacttt ataaagtttt tcatcatcac cacagcaatc acaaagagaa taattatgaa 3780
 tatacgcaag aggaaatgag aagggaatcc aaatgtcatt aaaaaaaaaa 3829

<210> 808

<211> 781

<212> DNA

<213> Homo sapiens

<400> 808

gcggcggagc tgtgagccgg cgactcgggt ccctgaggtc tggattcttt ctccgctact 60
 gagacacggc gggtaggtcc acaggcagat ccaactggga gttgaagtgt gaggtagagt 120
 gaagaggaac cagcaggctt ccggagggtt gtgtggtcag tgactcagag tgagaaggcc 180
 ctccaagtgc tgcctcctct catgcggtgc cagccccatg gaccttcttg tctcgtcacg 240
 gccataacta gggaggaagg agggccgagg agtgaggagg ctcaggcgaa gctgggggtgc 300
 tgttgggggt atccgagtcc cagaagcacc tggaaacccc acagaagatt ctggactccc 360
 cagacgggac caggagaggg acggcatgag cgacacacac aaacacagaa ccacacagcc 420
 agtcccagga gccagtaat ggagagcccc aaaaagaaga accagcagct gaaagtgcggg 480
 atctacacac tgggcagcag acagaagaag atcaggatac agctgagatc ccagtgcgcg 540
 acatggaagg tgatctgcaa gagctgcac agtcaaacac cggggataaa tctggatttg 600
 ggttccggcg tcaaggtgaa gataatacct aaagaggaac actgtaaaat gccagaagca 660
 ggtgaagagc aaccacaagt ttaaatgaag acaagctgaa acaacgcaag ctggttttat 720
 attagatatt tgacttaaac tatctcaata aagttttgca gctttcacca aaaaaaaaaa 780

a

781

<210> 809
 <211> 160
 <212> PRT
 <213> Homo sapiens

<400> 809
 Met Arg Cys His Ala His Gly Pro Ser Cys Leu Val Thr Ala Ile Thr
 5 10 15
 Arg Glu Glu Gly Gly Pro Arg Ser Gly Gly Ala Gln Ala Lys Leu Gly
 20 25 30
 Cys Cys Trp Gly Tyr Pro Ser Pro Arg Ser Thr Trp Asn Pro Asp Arg
 35 40 45
 Arg Phe Trp Thr Pro Gln Thr Gly Pro Gly Glu Gly Arg His Glu Arg
 50 55 60
 His Thr Gln Thr Gln Asn His Thr Ala Ser Pro Arg Ser Pro Val Met
 65 70 75 80
 Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu His
 85 90 95
 Leu Gly Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln Cys
 100 105 110
 Ala Thr Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro Gly
 115 120 125
 Ile Asn Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro Lys
 130 135 140
 Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln Val
 145 150 155 160

<210> 810
 <211> 624
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(624)
 <223> n=A,T,C or G

<400> 810
 atganaagga gatgacacaa aagtttagatc tcatcacaag tgatttggca gattaccagc 60
 agccctcat gatnggcacc gggacagtca cgaggaaggg ctccaccttc cggcccatgg 120
 acacggatgc cgaggaggca ggggtgagca ccatgcccgg cggccactat gactgcccgc 180

006280" E95T5960

```

agcgggcccgg ccgccacgag tacgcgctgc ccttggcgcc cccggagccc gactacgcca 240
cgcccatcgt ggagcggcac gtgctgcgcg cccacacgtt ctctgcgcag agcggctacc 300
gcgtcccagg gccccagccc ggccacaaac actccctctc ctcgggcggc ttctcccccg 360
tagcgggtgt gggcgcccag gacggagact atcaaaggcc acacagcgca cagcctgcgg 420
acaggggcta cgaccggccc aaagctgtca gcgccctcgc caccgaaagc ggacaccctg 480
actctcagaa gcccccaacg catcccggga caagtgcag ctattctgcc cccagagact 540
gcctcacacc cctcaaccag acggccatga ctgccctttt gtgaacacaa tgtgaaagaa 600
gcctgctgtg gtactgagcg tcgg                                     624

```

<210> 811

<211> 572

<212> DNA

<213> Homo sapiens

<400> 811

```

agcgggctgt gaggacgtc tgggccaggc tgcagcgca gcgttccgag ctgctgggct 60
ctttcgagga tgttctgata cgcgcgtcgg cctgcctgga ggaggcggcc cgggagcgcg 120
acggcctgga gcaggcgctg cggaggcgcg agagcgagca cgagaggag gtgcgcgctc 180
tgtacgagga gacggagcag cttcgggagc agagccggcg ccgcgcgagt cagaacttcg 240
cccgcgggga gcggagaagc cgtctggagc tggagctgca gatccgcgag caggaccttg 300
aacgcgcggg cctgcggcag cgggagttag agcagcagct gcacgcccag gctgcggagc 360
acctggaggc acaggcccag aactcccagc tgtggcgggc gcacgaggcg ctgcgaacgc 420
agctggaggg ggcgaggag cagatccgca ggctggagag cgaagcacga ggccgcccag 480
agcaaaccca acgagacgtg gtccgctct ccaggaacat gcagaaagag aaagtcagcc 540
tgctacggca actggagctg ctgaggagc tg                                     572

```

<210> 812

<211> 594

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(594)

<223> n=A,T,C or G

<400> 812

```

cggaagtgg cgcagcgcg ttgccaatgg tcgctccctg attnatgcc gctcgtgggtg 60
ttttgcgggc tgccgtacag cggcaagagc cggcgtgctg aagagttgcg cgtggcgctg 120
gctgccgagg gccgcgcggt gtacgtgggtg gacgacgcag ctgtcctggg cgcagaggac 180
ccagcggtgt acggcgattc tgcccgtgag aaggcattgc gtggagctct gcgagcctcc 240
gtggaacggc goctgagtcg ccacgacgtg gtcacctgg actcgcttaa ctacatcaaa 300
ggtttccggt acgagctcta ctgacctggca cgggcggcgc gcaccccgct ctgcctgggtc 360
tactgcgtac ggcccggcgg cccgatcgcg ggacctcagg tggcgggcgc gaacgagaac 420
cctggccgga acgtcagtg gagttggcgg ccacgcgctg aggaggacgg gagagcccag 480
gcggcgggga gcagcgtcct cagggaaact catactgcgg actctgtagt aaatggaagt 540
gcccgaggcc acgtacccaa ggaactggag cgagaagaat ccggggctgc ggag                                     594

```

<210> 813

<211> 561

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(561)
<223> n=A,T,C or G

<400> 813
tctgacacac gagaccgggt atcccatctc cgcgcccctc tgtgggtatt acacagccac 60
tagatgaagc caaacattgt tggaggtagt gaaatcttag actccaccat gtgtccagga 120
nccattgac gtctctctct ctgaaaactc cgtgtggccc tcgctctgca ctgtcatgag 180
gcggtgatgg agctagatac ccaccacgga caatgatcat cagtttgggg ttctctgggt 240
ctcacaggga cgcacattct aggggtagca cgacactccc cctgtagttg ctccacacaa 300
acgggatctc tcatccaggc gatacgtctg gtctgtggc atgtggctct cnacgaaaca 360
ccagggangc attatgttgg ggacttcttg gggctctgct ggtctctgct ccagacacga 420
ttaatccgaa atgtgtttaan tcgancacat ggggtccacgt ccaggacagc tcccatcgaa 480
ctctcnaggc tctctanctc agggatgaag gaggtnaagt gatcgatnct cacaagcgan 540
agctctcgcn cnatatctgc g 561

<210> 814
<211> 307
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(307)
<223> n=A,T,C or G

<400> 814
cntcngngng ttggttgtgt gggntnttct cgggtgattg ggtgmnatta ctggacccaa 60
ccnncgtgga aanggtggg nncgcggcgc ntctngcaga agtatcccga tttttttttt 120
tttttttttt tttttggngg agggaaantt ncagacatag ctttattgct gactcctgcc 180
cccttcanag ccctagtcac aggcnnacagg gntgttttgt aanttaaant ttcnggaaaa 240
tngngtntt tntgcatnca anagaagggn tgccaaangn ggggtattgc ttctgggtgg 300
nttacc 307

<210> 815
<211> 784
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(784)
<223> n=A,T,C or G

<400> 815
ggcagcagat ataatcagac tcttactcct gtacttctag aaatgatgca aacacttcaa 60
ggaccacaaa atgtggaaga tatgaatgca ctgttaatca aagatgctgt gtataatgct 120
gttggtatga gctgcttatg agctctttga cagtgttgat tttgatcagt ggtttaaaaa 180
ccagcttctt ccagaattac aagtcattca caataggtat aagccattgc gacgcagggt 240
gatttggtct atcggtcagt ggatttctgt gaaattcaag tctgacttaa gacccatgct 300

```

ttatgaagca atctgtaact tgcttcaaga tcaagattta gtggccgtat tgaaacagct 360
acaactttga agttaactgt tgatgatttt gaatttagaa cagatcagtt tctaccgtat 420
ttggaaacca tgttcacact acttttttcag ttactgcagc aagttacaga atgtgacaca 480
aagatgcatg ttttgcattgt cctttcttgt gtgatcgaaa gagtcaacat gcagatacga 540
ccatatgtgg gatgtttggt acaatatattg cccctccttt ggaagcagaa gtgaanaaca 600
caatatgttg agatgtgcta ttttgaccac acttattcat cttggtcagg gattangagc 660
agacagcaag acctgtccct ttcctgctcc agttattcac tgagtaccag atgtttcaca 720
gccttcncat gtttattttt ctggaaaatg ggtaaaaaat atnggtanga acctttggga 780
aaac 784

```

```

<210> 816
<211> 813
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1)...(813)
<223> n=A,T,C or G

```

```

<400> 816
ggcacgagca ggctgggaag aagtccttgc ttctcaaggc cacgtaccgg ccgcgtcctt 60
ccacccttgc cctttaaacc acagatgcc aatgatacgc caacagacac tacattcccc 120
agcagctgct gccagagccc tctttagct tctttatttt ctgtttcttt ccagctttcc 180
taccctccta tcccccttg tgtttgggccc acaattttga aataattttt attataggta 240
tgtgtgcgca aagccagatt tttataaggt aaaataaatt aagaatttaa acagtaaaag 300
ccagtgtctc aaaatgtcag cattaataatg tgaaggggac agcaggggtgt gaaccggaaa 360
cacacattgc caaacagttg ccaactgaac tgctgcttct catgggtcgt tcttttcttt 420
gcccttaagg tcaatgccag tgtccagacg agcagtgtag aaaagctccc tgtgtgggtt 480
gtcgtgaggt ctgcttgtat ctcttctact gcgttagttt cattagctct ttattctcct 540
tacgttcgag tgaatctgcc aagaacactg gtggatagta ttatcctaac acttttggtt 600
tgggggctgg gagggggcag ggaatagtga gctggcttta ccaccttcag gatctcgaat 660
tgggcgcttg aacctaaaga agattgtgga cttatcaaaa gtcaccgctc agtggttcgtc 720
aagcatgtat ttatgtgacn atcatactag ggaggggatg gttgggaatt cttccatgtg 780
caaatttngn cccgcaanaa gcaaaactgg ngt 813

```

```

<210> 817
<211> 229
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1)...(229)
<223> n=A,T,C or G

```

```

<400> 817
gaaactttta cattaatgat ttattaaaaa aaacaactcc ttgtcccact ccactgngct 60
gcttgtaatc tccatacatg gcctccattt tcaactgttt tnttggtcac anagctccaa 120
acanacacat ttttttttcc aggtaaaagc tgtttttagt ttgtagtaca aatgtgactg 180
catccaatac tgacacattg ttcctttggc ccacagtccc antcaccac 229

```

```

<210> 818
<211> 781

```

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(781)
<223> n=A,T,C or G

<400> 818
ggcacgaggt gtgtgtgtgt gtgtgtgtgt aacacatggg cattgggcct tccaggacaa 60
cttggttagg gctccagggt ggctctcag gcaggaacag gcttttttcc tcctgtcttt 120
tcctcacatc acgtcctgcc ccaggtcact gcataaataa gtgctttgga aagtattcat 180
ctagaaagta acataaatac tgtacataga aaagggttgc cgcccttag ccttcgcact 240
gccccagaga gctctccaca tattgcacac ggctcccca gccctgtggg gtccaggcct 300
ggctgtgtct ttggtagaag cttcagggac agttcctggg cagccccac atctncaccc 360
tgctcccaaa ggggagctct agggtagtca gtgggtacca gaagccttgc tcggcctcgc 420
tggtggcctt ctaccangga tgctttcaca aggatgagac agaatcccaa tggatgccc 480
ctgcttggac actctgctca aggtctgcat gtggcctggg aggagacagg caggctgang 540
gcaggtggac aggtgantcc tggccacana aggcaggctc acacccttca cangaatagg 600
tggtttgngc tgtcatctcg gcccacggtc tcctnntgcg ccaccccccc ttnttgaatc 660
gnaantcttc aaanccctta ccaccacttg atgaccnanc atttttangg cctggcttga 720
aggngggggc cttnggcccc ccnaaggggg aaatncccc ggngaatnc ccaangggga 780
a 781

<210> 819
<211> 199
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(199)
<223> n=A,T,C or G

<400> 819
cnnngtgga anggctgggn nngcgccgt tttcgnngta gtatcgcnt tttttttttt 60
tttttgtggg aggttntgcn gtntttgntt gctctctcaa attccaggaa ttgacttatt 120
taattaatgc ctgcaacctg tgctagcaaa tatttgnaca aaacnanttg tgttggngat 180
gttcttttgg gtcgggcag 199

<210> 820
<211> 211
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(211)
<223> n=A,T,C or G

<400> 820
nnnggcacga ggagagagag agagagagag agagagagag agagagagag agagagagag 60
agagagagag agagagagag agagagagag agagagagag agagagagag agagagagag 120
agacagtnc ntgtgtgtct ctctgtctcn aagtacncnc tgaggnatct gntntctgtn 180

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(264)
<223> n=A,T,C or G

<400> 823
ntcnatncct actangncaa actgactcgc ccctnagnca cctngtggtc canggctgcg 60
gagctgcgat acagccttcc ggggtctggn tggaaccccg acctntctg gtgtntntcc 120
ntccnccncc ccaacccgcc aagggcctgc ctttctnct gggcctttgc cagcgnnngg 180
ccanaccggg gccaaaccgg nccccgggca cattttaacc nagggcncnc ttntagaana 240
aaaccccggn tgatgttata aagg 264

<210> 824
<211> 520
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(520)
<223> n=A,T,C or G

<400> 824
tcaagcngcc cccantntga tggatatctg caaaattcnc cctttcacgc gccgcccgc 60
gcatgtctta ttatacaaca natccaactt ccctaagnng ntcacacatn ntaagggtatt 120
gttaacaaaa taggaaantc tattngaact aacaatcatc tctttgaatc tgcntatccc 180
attaaaagca ttttctctca tattctctat atcggttatg gncaatggat acccatctga 240
gctgggttgan ccttttaaat tnattatact taactttttg aaggctgtta taccacaggg 300
acaaacctaa ncaaccanca gatatacttg anggtntctc ctgtnatttc tcagattcca 360
atataccatt ttgccttnac acctacagcc cttaggggca tctctnttcc ncanaacaaa 420
ncattntcac taagacagnc tggggtntntn caccaatggc taccaaacct ctgnccgcna 480
cccaccgcnt aaanggcnga aattncnanc ccacacgggt 520

<210> 825
<211> 2064
<212> DNA
<213> Homo sapiens

<400> 825
cggtgcgctg agcgccggag gagcgtaggc agggcagcgc tggcgccagt ggcgacagga 60
gccgcgcgac cggcaaaaaat acacgggagg ccgtcgccga aaagagtccg cggtcctctc 120
tcgtaaacac actctctctc accggcgccct cccctctcgc tctgcgcgcc gcccggtggt 180
gcgcccagag cgcctccgac tgctatgtga ccgcgaggct gcgggaggaa ggggacaggg 240
aagaagaggc tctcccgcgg gagcccttga ggaccaagtt tgcggccact tctgcaggcg 300
tcccttctta gctctgcgcc gccctttctc gcagcctagg cggcccgggt tctcttctct 360
tctctgcgcg cccagccgcc tcgggtcccc gcgaccatgg tgacgatgga ggagctgcgg 420
gagatggact gcagtgtgct caaaaggctg atgaaccggg acgagaatgg cggcggcgcg 480
ggcggcagcg gcagccacgg caccctgggg ctgccgagcg gcggcaagt cctgctgctg 540
gactgcagac cgttctctgg gcacagcgcg ggctacatcc taggttcggt caacgtgcgc 600
tgtaaacacca tcgtgcggcg gcgggctaag ggctccgtga gcctggagca gatcctgccc 660
gccgaggagg aggtacgcgc ccgcttgccg tccggcctct actcggcggt catcgtctac 720

gacgagcgca gcccgcgcg cagagagcctc cgcgaggaca gcaccgtgtc gctggtggtg 780
 caggcgctgc gccgcaacgc cgagcgccacc gacatctgcc tgctcaaagg cggctatgag 840
 aggttttccct ccgagtaccc agaattctgt tctaaaacca aggccctggc agccatccca 900
 cccccggttc cccccagtgc cacagagccc ttggacctgg gctgcagctc ctgtgggacc 960
 ccactacacg accagggggg tcctgtggag atccttccct tcctctacct cggcagtgcc 1020
 taccatgctg cccggagaga catgctggac gccctgggca tcacggctct gttgaatgtc 1080
 tcctcggaact gcccaaacca ctttgaagga cactatcagt acaagtgcac cccagtggaa 1140
 gataaccaca aggcgcacat cagctcctgg ttcatggaag ccatagagta catcgatgcc 1200
 gtgaaggact gccgtggggc cgtgctggtg cactgccagg cgggcatctc gcggtcggcc 1260
 accatctgcc tggcctacct gatgatgaag aaacgggtga ggctggagga ggccttcgag 1320
 ttcgttaagc agcgccgcag catcatctcg cccaacttca gcttcatggg gcagctgctg 1380
 cagttcgagt cccaggtgct ggccacgtcc tgtgctgcgg aggtgctag cccctcggga 1440
 cccctgcggg agcggggcaa gacccccgcc acccccacct cgcagttcgt cttcagcttt 1500
 cgggtctccg tgggcgtgca ctcgggcccc agcagcctgc cctacctgca cagccccatc 1560
 accacctctc ccagctgtta gagccgcccc gggggcccca gaaccagagc tggctccag 1620
 caagggtagg acgggcccga tcggggcaga aagttgggac tgagcagctg ggagcaggcg 1680
 accgagctcc tccccatca tttctccttg gccaacgacg aggcagcca gaatggcaat 1740
 aaggactccg aatacataat aaaagcaaac agaacaactc aacttagagc aataacggct 1800
 gccgcagcag ccagggaaga ccttggtttg gtttatgtgt cagtttctact tttccgatag 1860
 aaattttctta cctcatTTTT ttaagcagta aggcttgaag tgatgaaacc cacagatcct 1920
 agcaaagtgt cccaaccagc tttactaaag ggggaggaag ggagggcaaa gggatgagaa 1980
 gacaagtttc ccagaagtgc ctggttctgt gtacttgtcc ctttgttgtc gttgtttag 2040
 ttaaaggaat ttcatttttt aaaa 2064

<210> 826

<211> 2109

<212> DNA

<213> Homo sapiens

<400> 826

tggcgccagc ggcgacagga gccgcgcgac cggcaaaaat acacgggagg ccgtcgccga 60
 aaagagtccg cggtcctctc tcgtaaacac actctcctcc accggcgccct cccctccgc 120
 tctgcgcgac gcccggtggtg gcgcccagag ccgctccgac tgctatgtga ccgcgaggct 180
 gcgggaggaa ggggacaggg aagaagaggc tctcccgcgg gagcccttga ggaccaagt 240
 tgcgccact tctgcaggcg tcccttctta gctctgcct gccctttct gcagcctagg 300
 cggcccagggt tctcttctct tctctgcgag cccagccgcc tcggttcccg gcgaccatgg 360
 tgacgatgga ggagctgcgg gagatggact gcagtgtgct caaaaggctg atgaaccggg 420
 acgagaatgg cggcggcgcg ggcggcagcg gcagccacgg caccctgggg ctgccgagcg 480
 gcggcaagtg cctgctgctg gactgcagac cgttcctggc gcacagcgcg ggctacatcc 540
 taggttcggt caacgtgcgc tgtaaacacca tcgtgcggcg gcgggctaag ggctccgtga 600
 gcctggagca gatcctgcc gccgaggagg aggtacgcgc ccgcttgcgc tccggcctct 660
 actcggcggt catcgtctac gacgagcgca gccgcgcgc cgagagcctc cgcgaggaca 720
 gcaccgtgtc gctggtggtg caggcgctgc gccgcaacgc cgagcgacc gacatctgcc 780
 tgctcaaagg cggctatgag aggttttccct ccgagtacc agaattctgt tctaaaacca 840
 aggccctggc agccatccca cccccggttc cccccagcgc cacagagccc ttggacctgg 900
 gctgcagctc ctgtgggacc ccactacacg accagggggg tcctgtggag atccttccct 960
 tcctctacct cggcagtgcc taccatgctg cccggagaga catgctggac gccctgggca 1020
 tcacggctct gttgaatgtc tcctcgact gcccaaacca ctttgaagga cactatcagt 1080
 caaagtgcac ccagtgga gataaccaca aggcgcacat cagctcctgg ttcatggaag 1140
 ccatagagta catcgatgcc gtgaaggact gccgtggggc cgtgctggtg cactgccagg 1200
 cgggcatctc gcggtcggcc accatctgcc tggcctacct gatgatgaag aaacgggtga 1260
 ggctggagga ggccttcgag ttcgttaaagc agcgccgcag catcatctcg cccaacttca 1320
 gcttcatggg gcagctgctg cagttcgagt cccaggtgct ggccacgtcc tgtgctgcgg 1380
 aggtgctag cccctcggga cccctgcggg agcggggcaa gacccccgcc acccccacct 1440

Glu Pro Leu Asp Leu Gly Cys Ser Ser Cys Gly Thr Pro Leu His Asp
180 185 190

Gln Gly Gly Pro Val Glu Ile Leu Pro Phe Leu Tyr Leu Gly Ser Ala
195 200 205

Tyr His Ala Ala Arg Arg Asp Met Leu Asp Ala Leu Gly Ile Thr Ala
210 215 220

Leu Leu Asn Val Ser Ser Asp Cys Pro Asn His Phe Glu Gly His Tyr
225 230 235 240

Gln Tyr Lys Cys Ile Pro Val Glu Asp Asn His Lys Ala Asp Ile Ser
245 250 255

Ser Trp Phe Met Glu Ala Ile Glu Tyr Ile Asp Ala Val Lys Asp Cys
260 265 270

Arg Gly Arg Val Leu Val His Cys Gln Ala Gly Ile Ser Arg Ser Ala
275 280 285

Thr Ile Cys Leu Ala Tyr Leu Met Met Lys Lys Arg Val Arg Leu Glu
290 295 300

Glu Ala Phe Glu Phe Val Lys Gln Arg Arg Ser Ile Ile Ser Pro Asn
305 310 315 320

Phe Ser Phe Met Gly Gln Leu Leu Gln Phe Glu Ser Gln Val Leu Ala
325 330 335

Thr Ser Cys Ala Ala Glu Ala Ala Ser Pro Ser Gly Pro Leu Arg Glu
340 345 350

Arg Gly Lys Thr Pro Ala Thr Pro Thr Ser Gln Phe Val Phe Ser Phe
355 360 365

Pro Val Ser Val Gly Val His Ser Ala Pro Ser Ser Leu Pro Tyr Leu
370 375 380

His Ser Pro Ile Thr Thr Ser Pro Ser Cys
385 390

<210> 828
<211> 453
<212> DNA
<213> Homo sapien

<400> 828
ggatcattta attgcatact ctatgaccac gcacatgtaa agccccttct gcaaaagaga 60
cctaaaccag atgagaagta ttattcatcc agcatatggg gaccaacatg tgatggcctc 120
gatcggattg ttgagcgcgtg tgacctgcct gaaatgcatg tgggtgattg gatgctcttt 180
gaaaacatgg gcgcttacac tgttgetgct gcctctacgt tcaatggctt ccagaggccg 240
acgatctact atgtgatgtc agggcctgcg tggcaactca tgcagcaatt ccagaacccc 300

```
<210> 829
<211> 452
<212> DNA
<213> Homo sapien
```

```
<210> 830
<211> 450
<212> DNA
<213> Homo sapien
```

```
<210> 831
<211> 395
<212> DNA
<213> Homo sapien
```

```
<210> 832
<211> 291
<212> DNA
<213> Homo sapien
```

<400> 832

```

ctgactcttc catctgtgca ggttgactga ggtcattcct gagttgcagt atgttgagag      60
ggtaatatatt ctgtctttctc taactcccca tactcccttg tcttccactc tccacttagg      120
agttttttgt gagttatgtc cttgttgctt ttgcctcttt ttctttctag ccttgattgt      180
gccagaagac aatgtcccta ttcacacact ctttctgctt ttctgtgggc aggaacatgg      240
aagggtgtgt gatggacgtg gactgtgaga gcgtctaccc cactgtgtag g                291

```

<210> 833

<211> 491

<212> DNA

<213> Homo sapien

<400> 833

```

ctgtagcttc tgtgggactt ccactgctca ggcgtcaggc tcaggtagct gctggccgcg      60
tacttggtgt tgctttgttt ggagggtgtg gtggtctcca ctccgcctt gacggggctg      120
ctatctgcct tccaggccac tgtcacggct tccgggtaga agtcacttat gagacacacc      180
agtgtggcct tggttgcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg      240
gcagccttg gctgacctag gacggctcagc ttggtccctc cgccgaagac cacattattg      300
ccgtcccacg tctgacagta atagtcagcc tcatccatag cctgggtccc gctgatggtc      360
agagtggctg tgttcccaga gttggagcca gagaagcgct cagggatccc tgaagaccgc      420
ttattatctt gataaatgac taccacaggg gactggcctg gcttctgttg ataccaacaa      480
gcagatacct g                491

```

<210> 834

<211> 308

<212> DNA

<213> Homo sapien

<400> 834

```

ctggctcagg tccacgccgc ggtaggtgaa cttgcggaag gtccgcttct tcttctgctc      60
tacttctgcc gtgctggaga acatcgaact gaacaagaag agtatgtatt cccgtgtgcc      120
agagtgccag gtcaccacat actattatgt tgggttcgca tatttgatga tgcgtcgta      180
ccaggatgcc atccgggtct tcgccaacat cctcctctac atccagagga ccaagagcat      240
gttccagagg accacgtaca agtatgagat gattaacaag cagaatgagc agatgcatgc      300
gctgctgg                308

```

<210> 835

<211> 472

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (472)

<223> n = A,T,C or G

<400> 835

```

ctgacatggt aactgtgatg cataaaactc gatcttctga tggggagtaa gtgcagaagg      60
tagaaatctc cgccccgcgg gggcttatct gtactggtag ttcattgctgt ggtctgcgtt      120
tgtgccatag ccgccttggt aggactggta ggagctggga gggccactgt agttctggcc      180
ggaccccggg gagttgtagt tcgactgtga gtagectcct tgtttgctt ggtatgagga      240
ggcgccccca gaacctccgc cgtagcccc gtgtgaccct gggttgtagg atgccccgcc      300
tgagccgtag ctgttccgcg cgcttcggcc tccactacca ctgtagtga atttgctctc      360
gtagntgtag tcggatccgc ccccgcccc gggagagttg tngganttcg agtaggagta      420
gctgccttgt ccatggttat agcctttctg cttgcctgtt ggagggccat ag                472

```

<210> 836
 <211> 354
 <212> DNA
 <213> Homo sapien

<400> 836
 ccagtgcac cttcagatag acacatgggtg accagagccc gccaggcttc tgcagggtggc 60
 agtgtcgagc aagtgtgaaga tgtctgtggg aaggagaagc tcctgaaatg aacgttctgc 120
 aaacagaagg ctgaggggtc ttccaggcat gtccagtcac taggagctgc caccgggtggg 180
 cttgagtgcc aggctctagg ctttgtgcag aaagcaccgc gggcgggggg cggttaaggga 240
 gagcaaaatg ggtctctctc aactgcagtc agtgcctctg ggaacacggc ctcacagaca 300
 gcacatattc tacgtcacag ctctagggtt tcaaggactt agccatccga cagg 354

<210> 837
 <211> 318
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(318)
 <223> n = A,T,C or G

<400> 837
 ctgaaaatga aggtaattaa aaccatggag gcgatcagcg aggttctcca ggaccttagg 60
 tttgatgcgg aatctgccga gtgatggcgg ctccccaggg atgcgccgag ggagatggga 120
 aacggggcgg atggcgccca gccagccct aactgccagc cacattgaag cggacattgg 180
 caaccgggtc cccagccatg cgcagaaccg tgggtagcat gtgcttggtg gtgatgtcct 240
 gccacagac ctcagacggc acattgatgc agaagagcgt antcatgcgg tgcaggtagt 300
 tggggtctcc ggacatgg 318

<210> 838
 <211> 277
 <212> DNA
 <213> Homo sapien

<400> 838
 ctgcgcgtcg ccaaagtgc aggcgggtgc gcctccaagc tctctaagat ccgagtcgtc 60
 cggaaatcca ttgcccgtgt tctcacagtt attaaccaga ctcagaaaga aaacctcagg 120
 aaattctaca agggcaagaa gtacaagccc ctggacctgc ggcctaagaa ggcacgtgcc 180
 atgcgcgcc ggctcaacaa gcacgaggag aacctgaaga ccaagaagca gcacgggaag 240
 gagcggctgt acccgtgcg gaagtacgc gtcaagg 277

<210> 839
 <211> 276
 <212> DNA
 <213> Homo sapien

<400> 839
 ccaaggaatg caggctgtac tatctgcgaa atggagaacg tatttcagtg tcggcagcct 60
 ccaagctgct gtccaacatg atgtgccagt accggggcat gggcctctct atgggcagta 120
 tgatctgtgg ctgggataag aagggtcctg gactctacta cgtggatgaa catgggactc 180
 ggctctcagg aaatatgttc tccacgggta gtgggaacac ttatgcctac ggggtcatgg 240

acagtggcta tcggcctaata cttagccctg aagagg

276

<210> 840

<211> 453

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(453)

<223> n = A,T,C or G

<400> 840

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ccttctttgc | catgaccaag | ctctttcagt | ccaatgatcc | cacactccgt | cggatgtgct | 60 |
| acttgaccat | caaggagatg | tcttgcatg | cagaggatgt | catcattgtc | accagcagcc | 120 |
| taacaaaaga | catgactggg | aaagaagaca | actaccgggg | cccggccgtg | cgagccctct | 180 |
| gccagatcac | tgatagcacc | atgctgcagg | ctattgagcg | ctacatgaaa | caagccattg | 240 |
| tggacaaggt | gcccagtgtc | tccagctctg | ccctcgtgtc | ttccttgcac | ctgctgaagt | 300 |
| gcagctttga | cgtgggtcaag | cgctgggtga | atgaggctca | ggaggcagca | tccagtata | 360 |
| acatcatggt | ccagtaccac | gcactanggc | tcctgtacca | tgtgcgtaag | aatgaccgcc | 420 |
| tagccgtcaa | taagatgata | agcaagggtc | cac | | | 453 |

<210> 841

<211> 142

<212> DNA

<213> Homo sapien

<400> 841

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| agcctctcta | gtggcagagc | agctcacact | ccctccgctg | ggaacgatgg | cttctgccta | 60 |
| gtacctatcc | ttgtgtttct | gatgcagtgg | tagcattggt | tcaagttctc | tcctgctgtg | 120 |
| gtcagagtgt | cttcgatgtt | gg | | | | 142 |

<210> 842

<211> 83

<212> DNA

<213> Homo sapien

<400> 842

| | | | | | | |
|------------|------------|------------|------------|-----------|------------|----|
| cctaaaagca | gccaccaatt | aagaaagcgt | tcaagctcaa | caccactac | ctaaaaaatc | 60 |
| ccaaacatat | aactgaactc | ccc | | | | 83 |

<210> 843

<211> 482

<212> DNA

<213> Homo sapien

<400> 843

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ccatcgggtg | ctggcagatg | cggcacctca | agagcttctt | tgaagccaag | aagcttgtgt | 60 |
| agctgtccca | ggcgtcacia | cccacctcc | caggctgggg | gagaaaggac | ctcctggaac | 120 |
| tgacttcttc | tgtcaggagg | actggtttcc | agccatacct | gttctggaag | ggagaggggc | 180 |
| tggaggcacc | cacaggcaca | agctgaaggc | agcagcttgg | ctaatactga | gcaggtagtg | 240 |
| gggcaaattc | ctgcctcttc | tctctggcct | ctgggccgtt | tggtagtaat | caccagggg | 300 |
| ctggtaaagc | ccctcctctt | ggcacctcag | aatcacagtg | ttactgatca | gggatgtgag | 360 |
| gctgctgttg | gggggtggggg | gaggggaatg | ggcaggcaag | ccagtcttct | gtcttctctt | 420 |

gctaacttag ggttttgagc aggttggggg tatggtgcct gtcataccca cctgccaccc 480
tg 482

<210> 844
<211> 534
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(534)
<223> n = A,T,C or G

<400> 844
ccagatTTTT caagtttaaa ggaggaaact gcttattgga aggaactttc cttgaagtat 60
aagcaaagct tccaggaagc tcgggatgag ctagttgaat tccaggaagg aagcagagaa 120
ttagaagcag agttggaggc acaattagta caggctgaac aaagaaatag agacttgcag 180
gctgataacc aaagactgaa atatgaagcg gaggcattaa aggagaagct agagcatcaa 240
tatgcacaga gctataagca ggtctcagtg ttagaagatg atttaagtca gactcggggc 300
attaaggagc agttgcataa gtatgtgaga gagctggagc aggccaacga cgacctggag 360
cgagccaaaa gggcaacaat agtttctactg gaagactttt gaacaaaggc taaaccaggc 420
cattgaacga aatgcatttt tagaaagttg aacttgatga aaaaggaatc tttgttggtc 480
tctgtacaga ggttnaagga tgaagcanga gatttaaggc aagaactagc agtt 534

<210> 845
<211> 175
<212> DNA
<213> Homo sapien

<400> 845
tcgacctgtg gcaaagtgtg ctacctgcc aagcgcaaga gaaagtataa ctggagtgcc 60
aaggctaaaa gacgaaatac caccggaact ggtcggatga ggcacctaaa aattgtatac 120
cgcagattca ggcattgatt ccgtgaagga acaacaccta aaccaagag ggcag 175

<210> 846
<211> 179
<212> DNA
<213> Homo sapien

<400> 846
cgcgtggaca gttgagaggg gtctgtgtga aggcacttgt cacgagcttc aatactgccg 60
ccgtcccagg atgggagaac tgcgcagcag gaagggcact tctgaaagca cagtggagag 120
atcgtctggag cgggcgttct gggcaggagg aagcacagac ggcaggcagg gtggactgg 179

<210> 847
<211> 410
<212> DNA
<213> Homo sapien

<400> 847
ccacaaaaac cagtcacaag acctggagtt gtctgtgcag atgtacgcc aagccgccct 60
ggatggagac tcccaggat tttttaacct ggccttgcta atcgaggaag gtacgataat 120
cccacacat atcttgatt tcttggaat tgactcaact ctccattcta ataacatctc 180
cattctccag gaactgtacg aaaggtgctg gagccacagt aacgaggagt ccttcagccc 240


```
<210> 848
<211> 557
<212> DNA
<213> Homo sapien
```

| | | | | | | | | |
|-------------|-------------|-------------|------------|------------|-------------|--|--|-----|
| <400> | 848 | | | | | | | |
| cacgggcccc | cagccctgtg | tgggcottgt | ctgtctcagc | tcaaccacag | tctgacacca | | | 60 |
| gagcccactt | ccatcctctc | tgggtgtgagg | cacagcgagg | gcagcatctg | gaggagetct | | | 120 |
| gcagcctcca | cacctaccac | gacctcccag | ggctgggctc | aggaaaaaac | agccactgct | | | 180 |
| ttacaggaca | gggggttgaa | gctgagcccc | gcctcacacc | cacccccatg | cactcaaaga | | | 240 |
| ttggatttta | cagctacttg | caattcaaaa | ttcagaagaa | taaaaaatgg | gaacatacag | | | 300 |
| aactctaana | gatagacatc | agaaatttgt | aagttaagct | ttttcaaaaa | accagcaatt | | | 360 |
| ccccagcgta | gtcaagggtg | gacactgcac | gctctggcat | gatgggatgg | cgaccogggca | | | 420 |
| agcttttcct | ctcgcagatgc | tctgctgctt | gagagctatt | gctttgttaa | gatataaaaa | | | 480 |
| ggggtttctt | tttgtctttc | tgtaaaggngg | acttcagct | tttgattgaa | agtcctaggg | | | 540 |
| tgaattctatt | tctactctg | | | | | | | 557 |

| | | | | | | | |
|-------------|------------|------------|------------|------------|------------|--|-----|
| <400> | 849 | | | | | | |
| ctgatggtttt | ggaaatgaga | gaactacagt | ggtgaagaga | ccaggaggca | gctctcagt | | 60 |
| aaaccaacat | tgcggatgcc | cttcgtgagc | cttctcagtc | ccagcaggaa | gccacaaca | | 120 |
| ctggcctccc | cagcctgcct | gctgacaaca | cctaggctta | ctttatctaa | aatcagagt | | 180 |
| taccaggtct | gtagcagaaa | ataatcaact | aaatgtcagg | gacctatgag | tcatttaaaa | | 240 |
| caaaagagga | agtgaagcc | attaggcaag | ctatgtgctg | ggctgctaac | gtagccctg | | 300 |
| cagggagggg | tcaggagcgc | gctgcagtga | gccttgggtc | tcgcaggccc | agccctgctg | | 360 |
| caaggagcca | gggcacccag | gaaacatcag | cacacacaca | cacaggacc | ctcccttcac | | 420 |
| gtcacttggt | ttgctgccct | aaatggcttc | ttgcacccta | acccctgac | ctggaagaag | | 480 |
| gcagagagac | tggcccgtag | agagacctgc | aattctacgc | aaqct | | | 525 |

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| <400> 850 | | | | | | |
| cctcttgagg | cacatccttt | actgcattgt | ggacagcgag | tgtaagtcaa | gggatgtgct | 60 |
| ccagagttac | tttgacctcc | tgggggagct | gatgaagttc | aacgttgatg | cattcaagag | 120 |
| attcaataaa | tatatcaaca | cggatgcaaa | gttccaggta | ttcctgaagc | agatcaacag | 180 |
| ctccctggtg | gactccaaca | tgctggtgcg | ctgtgtcact | ctgtccctgg | accgatttga | 240 |
| aaaccaggtg | gatatgaaag | ttgccgaggt | actgtctgaa | tgccgcctgc | tcgcctacat | 300 |
| atcccaggtg | cccacgcaga | tgtccttcct | cttccgcctc | atcaacatca | tccacgtgca | 360 |

gacgcatcat aaatgggtttt ctttaagtga atggaagagt ttgacagaga tacacctttg 360
 taagaaaaca ttaagaatgc tggctgactg tgggtggctca cacctgtatt cccagcactt 420
 tgggagg 427

<210> 855
 <211> 311
 <212> DNA
 <213> Homo sapien

<400> 855
 ccagtattcc tggaggatat aacactgaca tcagcagggg tttcaatggc aacaattgca 60
 cgagctgccg gcagaagctt ctcccagggt ctcttgagat ttatgatata gatgccatca 120
 cttttccttt tatagatgta ctgttccatc tgggaagtcaa gattgggtgcc acctaagtgg 180
 gttcctgctg caaggaactt aaggacatcc tcctccttca tttgcaggac atcaagggct 240
 ccggacattg tgaaagtttc cctttaagtt acgacgggaa tccagaacaa cgccgtatgg 300
 acccctctgc a 311

<210> 856
 <211> 328
 <212> DNA
 <213> Homo sapien

<400> 856
 cctatggaag tttgggtgctt tgctccctgt gtttgcgaaa caggtatctc gtgatttcag 60
 aaaagcttga ggagattaag tctttccggg agctgacctg cctggatctt tcctgttgca 120
 agcttgagaga tgagcatgaa cttctagaac atctcaccaa tgaagccctg tctagtgtaa 180
 ctgagctcca cctgaaggat aattgtctat ctgatgctgg ggtgcggaag atgacagcac 240
 cagttcgagt gatgaaaaga ggtatccaat gcctgcatct gtgatctcag ggttacatga 300
 taagtctaata aatgttagat tctcaagg 328

<210> 857
 <211> 502
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(502)
 <223> n = A,T,C or G

<400> 857
 ctgaccggac cggctcatgcc cgtccggaac gtctataaga aggagaaagc tcgagtcac 60
 actgaggaag agaagaattt caaagccttc gctagtctcc gtatggcccg tgccaacgcc 120
 cggctcttcg gcatacgggc aaaaagagcc aaggaagccg cagaacagga tgttgaaaag 180
 aaaaaataaa gccctcctgg ggacttgga tcaagtcggca gtcattgctgg gtctccacgt 240
 ggtgtgtttc gtgggaacaa ctgggcctgg gatggggctt cactgctgtg acttcctcct 300
 gccaggggat ttggggcttt cttgaaagac agtccaagcc ctggataatg ctttaactttc 360
 tgtgttgaag cactgttggg tgtttgggta gtgactgatg taaaacgggt ttcttgtggg 420
 gaggttacag aggctgactt cagagtggac ttgtgttttt tcttttttaa gangtaaggt 480
 tgggctgggtg ctcacagacc tc 502

<210> 858
 <211> 411
 <212> DNA

<213> Homo sapien

<400> 858

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cgcccgaggt | ccttaatagt | taagttacag | ctaagaatgt | catgtcttgg | gttggaattt | 60 |
| tcatttttag | caccgttaat | gtattcaact | aaatctatgt | tagcaccttg | tctccaggca | 120 |
| gaacaacaaa | ccatccaaac | attttaaaca | ttgggggaaa | cacgaagggg | agggttaaag | 180 |
| acagaatcca | gtactgtgga | aggagtggat | ttagatcaca | agatccttgt | cgatatcctt | 240 |
| ctgcttgatg | ccgaagcagc | cggcccactc | atccagggcg | atgtacttgt | cattgtccag | 300 |
| gtcacaggtc | tcgaaaaagc | gggtggtgca | atgctccatg | gggatgaggg | gagcacgcag | 360 |
| tggagccagc | tcggtgtggg | agaggtaccc | gtcaatgggg | tgctggtcca | g | 411 |

<210> 859

<211> 232

<212> DNA

<213> Homo sapien

<400> 859

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| aaatcacaga | gggacttagt | attccattaa | tgcaaattgga | aacattaagt | tcatcatcag | 60 |
| atgataaaag | gaaaaaaaaa | acctgatact | catctcaaaa | gacgcagaga | agacatctgc | 120 |
| ataaatccag | tacctattat | tatttcaaat | ttaaaaactt | cttctttttt | aagagatagg | 180 |
| gtatcactat | gttgcccagg | ctgatcttga | actcttggcc | tcagatgatc | ct | 232 |

<210> 860

<211> 235

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(235)

<223> n = A,T,C or G

<400> 860

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tgcccagaaa | ggaaggggct | attgcctcct | cccagccacg | ttccctttcc | tcctctccct | 60 |
| cctgtggatt | ctcccatcag | ccatctgggt | ctcctcttaa | ggccagttga | agatgggtcc | 120 |
| ttacagcttc | ccaagttagg | ttagtgatgt | gaaatgctcc | tgtccctggc | cctacctcct | 180 |
| tcctgtcccc | cacccttgca | taaggcagtt | gttggttttc | ttccccaatn | ctttt | 235 |

<210> 861

<211> 457

<212> DNA

<213> Homo sapien

<400> 861

| | | | | | | |
|-------------|------------|------------|-------------|-------------|-------------|-----|
| ccaaaggaaa | gttggaaggc | aactgacaga | ttctgccttt | taggtacttg | aactggcagg | 60 |
| aaatgcatca | aaagacttaa | aggtaaagcg | tattaccctt | cgtcacttgc | aacttgctat | 120 |
| tcgtggagat | gaagaattgg | attctctcat | caaggctaca | attgctgggtg | gtggatatgtt | 180 |
| aactttctaac | attttaaaaa | atttcttcag | aggaagggaat | tttttgctgc | ttttaattag | 240 |
| tttttccagg | agaggaaatt | taagtatat | ttcaatgatg | gaagtatggg | tgtatcatga | 300 |
| aatttgatgt | atatgtataa | ctcaatgaat | ttttacctca | tacttgagct | gcatgttttt | 360 |
| aaagatacct | ttcaagttga | acagtataca | ctttcttggg | ttcaaatact | gtgatttttt | 420 |
| aaaaaatctt | aagtagaatt | aattcctgtc | actcccc | | | 457 |

<210> 862

<211> 561
 <212> DNA
 <213> Homo sapien

<400> 862
 ccaggtcatc accattggca atgagcgggt ccggtgtccg gaggcgctgt tccagccttc 60
 cttcctgggt atggaatctt gcggcatcca cgagaccacc ttcaactcca tcatgaagtg 120
 tgacgtggac atccgcaaag acctgtacgc caacacgggt ctgtcgggcg gcaccaccat 180
 gtatccgggc attgccgaca ggatgcagaa ggagatcacc gccctggcgc ccagcaccat 240
 gaagatcaag atcatcgcac ccccagagcg caagtactcg gtgtggatcg gtggctccat 300
 cctggcctca ctgtccacct tccagcagat gtggattagc aagcaggagt acgacgagtc 360
 gggcccctcc atcgtccacc gcaaattgctt ctaaaccggac tcagcagatg cgtagcattt 420
 gctgcatggg ttaattgaga atagaaattt gccctgggca aatgcacaca cctcatgcta 480
 gcctcacgaa actggaataa gccctcgaaa agaaattgtc cttgaagctt gtatctgata 540
 tcagcactgg attgtagaac t 561

<210> 863
 <211> 291
 <212> DNA
 <213> Homo sapien

<400> 863
 ccatagctgt cccacctatg gttttaaaaa cagactgtaa cttgatcttc tgaaatcctt 60
 ctogaaccac aactcgttct gttaaagaaa tcctaggaaa gaagtcctac tgatattgtc 120
 gatagtctcc aaaagggtgag gaaggtaact gagttgaagg caactgggag gggctctctg 180
 caaactgagg accattggaa aactgtgcag aggcaaattc tgtcaacaag ataccagctc 240
 cttcaattaa agctaggaga atgccacca ttgcggctga cccaaccatg g 291

<210> 864
 <211> 265
 <212> DNA
 <213> Homo sapien

<400> 864
 ctgaactttt ccacctggag tccttgaggaa taccggacgt gatcttcttt tataggtcca 60
 atgatgtgac ccagtcctgc agttctggga gatcaaccac catccgcgtc aggtgcagtc 120
 cacagaaaac tgtccctgga ggtttgctgc tgccaggaac gtgctcagat gggacctgtg 180
 atggctgcaa cttccacttc ctgtgggaga gcgcggctgc ttgcccgtc tgctcagtgg 240
 ctgactacca tgctatcgtc agcag 265

<210> 865
 <211> 144
 <212> DNA
 <213> Homo sapien

<400> 865
 cctccacctg cgttttgatc tagatgagca tattgtccat ctcccacagc ttgtccgggt 60
 tccgcaggta cggccgcccg tgctcgcgcg tcagcgacgc gatgtcctcg cgcacatcgt 120
 tgatgaccgg gagcagaaac tgct 144

<210> 866
 <211> 241
 <212> DNA
 <213> Homo sapien

<400> 866
 ctggctgtaa gtagcttcat agcaccagtc tttgagaatg tcaagctctc cagaaatcat 60
 ggctccagg acattgggga tgatgtcggt ctgcactgt ttcagaaacc ggtccttgtc 120
 aaaggccggg tccaccggga ggatctcgt gagcacctcc gacatctctg tcttgagaa 180
 caggccccc agcaagtcgg tgacctgtc cgtaaggggc cgggatgcc ggatgaacgc 240
 g 241

<210> 867
 <211> 364
 <212> DNA
 <213> Homo sapien

<400> 867
 cctgggcccg ctgacttcag ggtgaggcca cagctactgc agcgcttttt atttatttat 60
 ttatttactg agatggagtc ttgctctgtc acccaggctg gagtgcagtg gtgcaatctc 120
 ggctcactgc aacctctgcc tcctgggctg cagtgtattct cctgcgttca agtaattctc 180
 ctgcctcggc cttctgagta gttgggatta caggcatatg ccaccacact tggctaattt 240
 ttcgatattt tagtagaaat ggggtttcac catgttggcg aggctgggtc cgaactcctg 300
 acctcaagga tcctcctgcc tcggcctcct aagggtgctgg gattgcagggt gtgagccacc 360
 acgt 364

<210> 868
 <211> 472
 <212> DNA
 <213> Homo sapien

<400> 868
 ccaccagtcc acagatgtga ctggtaaggg atctagtaac agaggatgga gttgggcaga 60
 atattatcct ggatgatatg caccagcac taggatacac ctttcattag aatgaagaga 120
 acagacaaag cctcagaaa agatacaaag gcagagacat tgattagaac attatctcat 180
 aacagagggtg gggccattac ccaccattat tgtaaaataa ctgtaactaa ccaaaacaca 240
 tacaggcttc tttaatggag ttaataaaaac tatggcacat tgggaatcag gggcagaggt 300
 actgtttcca gacggaaaac tgggataaag ggagccatgc tgacagggcc ttattccagt 360
 ctaggttgtt agaaaggagc cctagcccag aaatgacagc aaatagccat aatcattatg 420
 tggggctgaa ccagaggaag ccaggctgag ccaagaagct ggaagtatct tg 472

<210> 869
 <211> 368
 <212> DNA
 <213> Homo sapien

<400> 869
 cctttcttgt aagtgaagaa aaaggaatgc agcaaagaag agttcgacat tggagtcctt 60
 agttccatca ggatccatt cgcagccttt agcatcatgt agaagcaaac tgcacctatg 120
 gctgagatag gtgcaatgac ctacaagatt ttgtgttttc tagctgtcca ggaaaagcca 180
 tcttcagtct tgctgacagt caaagagcaa gtgaaaccat ttccagccta aactacataa 240
 aagcagccga accaatgatt aaagacctct aaggctccat aatcatcatt aaatatgccc 300
 aaactcattg tgacttttta ttttatatac aggattaaaa tcaacattaa atcatcttat 360
 ttacatgg 368

<210> 870
 <211> 411
 <212> DNA

<213> Homo sapien

<400> 870

```

ggcgtgtcct tggacttaga gagtggggac gtcgggcttc ggagcgggag tgttcgttgt      60
gccagcgact aaaaagagaa ttaaatatgg gtgatgttga gaaaggcaag aagattttta      120
ttatgaagtg ttcccagtgc cacaccgttg aaaagggagg caagcacaag actgggccaa      180
atctccatgg tctcttttggg cgggagacag gtcaggcccc tggatactct tacacagccg      240
ccaataagaa caaaggcatc atctggggag aggatacact gatggagtat ttggagaatc      300
ccaagaagta catccctgga acaaaaatga tctttgtcgg cattaagaag aaggaagaaa      360
gggcagactt aatagcttat ctcaaaaaag ctactaatga gtaataattg g              411

```

<210> 871

<211> 385

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(385)

<223> n = A,T,C or G

<400> 871

```

tttttttttt ttnnnttttt ttttttnaaa gattcacttt atttattcat tctcctccaa      60
cattagcata attaaagcca aggaggagga gggggggtga ggtgaaanat ganctggagg      120
accgcaatag gggtaggtcc cctgtggaaa aagggtcana ggccaaagga tgggaggggg      180
tcaggctgga actgagganc aggtgggggc acttntccct ntaacactnt cccctgttga      240
agctntttgt gacgggcnan ctcaggccct gatgggngac ttencaggcg tanactttgt      300
gtttctcgna ntctgctttg ctcanctgca ggggtgctgnt gaggctgtan ggtgctgtcc      360
ttgctgtcct gctntgngac actct              385

```

<210> 872

<211> 184

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(184)

<223> n = A,T,C or G

<400> 872

```

cttccttcgg tctttantat ttttgattgt tatgtaaaac tcgcttttat tttaatattg      60
atgtcagtat ttcaactgct gtaaaattat aaacttttat acttgggtaa gtccccagg      120
ggcgagtcc tcgctctggg atgcaggcat gcttctcacc gtgcagagct gcacttggcc      180
tcag              184

```

<210> 873

<211> 397

<212> DNA

<213> Homo sapien

<400> 873

```

ctgtgggctc tgaatggcgt ccctttggct atccacgccg ccggcgacca ctgaattctg      60
tggttctaca acagggctct gctgaccgaa ttgtcagaga cgtccaggaa ttcacgata      120

```

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| accccaagtg | gtacactgac | agaggcattc | cttacagacg | tggctacctg | ctttatgggc | 180 |
| ccccagggtg | cggaaagagc | agttttatca | cagccctggc | tggggaactg | gagcacagca | 240 |
| tctgcctgct | gagcctcacg | gactccagcc | tctctgatga | cggactcaac | cacctgctga | 300 |
| gcgtggcccc | gcagcagagc | ctggtaactcc | tggaggatgt | ggatgctgct | tttctcagtc | 360 |
| gagacttggc | tgtggagaac | ccagtaaagt | accaagg | | | 397 |

<210> 874

<211> 156

<212> DNA

<213> Homo sapien

<400> 874

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccagaagaac | actatgccat | ggttgcactg | aattttgtgc | ctactctagg | gcaaacagaa | 60 |
| ttacaatcga | aggagttcct | atctatctgt | aaagaagaga | acatgaaatt | ctgttggcag | 120 |
| aagcagcatt | ttgaagaaat | aaaaggttca | ctgcag | | | 156 |

<210> 875

<211> 512

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(512)

<223> n = A,T,C or G

<400> 875

| | | | | | | |
|------------|------------|-------------|------------|------------|-------------|-----|
| ccagcatagc | gaaaacttgt | ctctactaaa | aatacaaaaa | ttagtcaggc | atggtgggtgc | 60 |
| acgtctgtaa | taccagcttc | tcaggaggct | gaggcacgag | gatcacttga | accaggagg | 120 |
| aggaggttgc | agtgagctga | gatcatgcc | gggcaacaga | atgagacttt | gtttaaaaaa | 180 |
| aaaaaaagt | acttgattta | agggaaaaaa | tgactggcta | tattcagtca | gatatggcaa | 240 |
| agagtctcaa | ggtgttaatg | tgaatgatta | aggtcttggg | gggggtgtcc | cctatcagac | 300 |
| tacagggtgt | tagaggcaca | gaaaaagggtg | cagttgggtt | cttaatgtga | aatgatgaga | 360 |
| agcacaactc | cagtgtgtct | ctttgtgtag | aatgtcagca | gacaccccct | gctagatgtg | 420 |
| ctggatcatg | ggaaagcatt | tccatttgtt | aatagattgt | tcagaagttt | taatttatga | 480 |
| tgggtgtggt | ggctcatgcc | tgtngtccca | gc | | | 512 |

<210> 876

<211> 199

<212> DNA

<213> Homo sapien

<400> 876

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| cctgtgccgg | gccccagggc | tggcagccac | cagctcctct | tccaggcatg | ggggacaccc | 60 |
| tgacaggatc | cggaaagtctc | catttaccca | aaaatgcaag | agccatgatc | agtcattggcg | 120 |
| acactgcagg | cgggtactgag | tgaccatgtc | cagtccggct | ccgtccctcc | cacacggggg | 180 |
| acaagcttct | ccgaggagg | | | | | 199 |

<210> 877

<211> 486

<212> DNA

<213> Homo sapien

<400> 877


```
<210> 878
<211> 363
<212> DNA
<213> Homo sapien
```

```
<210> 879
<211> 365
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|------------|------------|------------|------------|--|-----|
| <400> | 879 | | | | | | |
| gcccattgcc | gcgtgtggtc | agcacgcaca | acttgtggct | gctgtccttc | ctgaggaggt | | 60 |
| ggaatgggag | cacagccatc | acagacgata | ccctgggttg | cactctcacc | attacgctgc | | 120 |
| ggaatctaca | accccatgat | gcgggtctct | accagtgcc | gagcctccat | ggcagtggag | | 180 |
| ctgacacctt | caggaaggtc | ctggtggagg | tgctggcgag | ccccctggat | caccggaatg | | 240 |
| ctggagatct | ctggttcccc | ggggagctct | agagcttcga | ggatgcccat | atggagcaca | | 300 |
| gcatctccag | gagcctcttg | gaaggagaaa | tcccttccc | accacttcc | atccttntcc | | 360 |
| tctctg | | | | | | | 365 |

| | | | | | | | |
|-------------|------------|-------------|------------|------------|-------------|--|-----|
| <400> | 880 | | | | | | |
| ccatctcccc | tcaccccaac | ctggataaaa | tgttacacta | cccactaata | taaccactga | | 60 |
| cacacaaaac | aagctccttc | cagttttaaca | ttgaacatca | atctacattt | ccagtgaatg | | 120 |
| agctaaaact | atgagcaggc | cattcaactt | ttcatgatac | atttagtgct | cagaaattggt | | 180 |
| tgtattccatt | agcctgcctt | atagctcagg | tggcccaaga | tggagcctat | catcttcctt | | 240 |
| qqqqtgtttg | gtgtttccaa | gtaggagcat | aaaaaggata | ccgtccccca | ccccaccacc | | 300 |

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccatcccaca | taccctcact | ggcatccagg | agaccagcag | cagggtcaag | accccaaagt | 360 |
| ttgggcacca | caaataatgt | gatatgtgcc | aggagcacgg | ggggtagggg | tgaaagagaa | 420 |
| aaacaataag | g | | | | | 431 |

<210> 881
 <211> 335
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| <400> 881 | | | | | | |
| ccacagaggt | ggtattacaa | aatatacaaa | gtgggtttctt | tctttacatt | tcatagaaga | 60 |
| agcctgcctc | atttccaaat | gagagcacta | gaagcacaaa | tcatgcagac | cattttactat | 120 |
| ataacttatg | aaaaatgctg | tacagggtcg | tgactataga | tatagagtat | ttgggtctgt | 180 |
| ttgggaattg | atatctacaa | gggggagggt | caggggagga | ctgtctgata | tcttgacttg | 240 |
| ctgggatggt | ggagaagctg | ggatggggga | ggccccaatc | ttgctgcacg | gctacaccca | 300 |
| ctcctccttt | cctagataag | gctggagcgc | actgg | | | 335 |

<210> 882
 <211> 353
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| <400> 882 | | | | | | |
| atgcactcaa | agattggatt | ttacagctac | ttgcaattca | aaattcagaa | gaataaaaaa | 60 |
| tgggaacata | cagaactcta | aaagatagac | atcagaaatt | gttaagttaa | gctttttcaa | 120 |
| aaaatcagca | attccccagc | gtagtcaagg | gtggacactg | cacgctctgg | catgatggga | 180 |
| tggcgaccgg | gcaagctttc | ttcctcgaga | tgctctgctg | cttgagagct | attgctttgt | 240 |
| taagatataa | aaaggggttt | ctttttgtct | ttctgtaagg | tggacttcca | gctttttgatt | 300 |
| gaaagtccta | gggtgattct | atttctgctg | tgatttatct | gctgaaagct | cag | 353 |

<210> 883
 <211> 193
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| <400> 883 | | | | | | |
| ctggcagaga | agaatggcta | cgtgactgtc | agtgagatca | aagccagtct | taaatgggag | 60 |
| accgagcgag | cgcggaaggt | gccggaacac | ctgctgaagg | aaggggttggc | gtggctggac | 120 |
| ttacaggccc | caggggaggc | ccactactgg | ctgccagctc | tcctcactga | cctctactcc | 180 |
| caggagatta | cag | | | | | 193 |

<210> 884
 <211> 461
 <212> DNA
 <213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| <400> 884 | | | | | | |
| ctgaagaacc | ccatcagcgg | gctgttagaa | tatgccagct | tcgctagtca | aacctgtgag | 60 |
| ttcaacatga | tagagcagag | tggaccaccc | catgaacctc | ggtaagagac | caccagggaa | 120 |
| ctgtacctag | ggttggggtc | aggtgctttt | gctcctgacg | cagtcttggc | tgattttgtga | 180 |
| gcagtgtgtg | ttggtggcgc | ctatcttttc | ctccttccct | tctgcctttt | agctaaattc | 240 |
| cccttgattg | gccctttctc | cagatattga | gcagggaata | tagaccttgg | accagccaga | 300 |
| atcttggtctg | aacaaggggg | aggttgactc | tgttggctgt | aatgaagctt | ctttagaaat | 360 |
| gattggtttt | ggcgtacgc | ggtggctcat | gcctgtaatc | ccagcacttt | ttgaggccga | 420 |

461

<211> 266

<212> DNA

<213> Homo sapien

 $\langle 220 \rangle$

<221> misc feature

 $\langle 222 \rangle \quad (1) \dots (266)$

<223> n = A, T, C or G

<400> 885

ctgcgaatgct tcancacact tcagcaccga ggctgggcat gaggggtccg tcaccaccac 60

atcaaataacc cctaaagcaa tatctttgtt atgggcactt gaatggtgct gcttcacaga 120

ggctgcacca ccagtcacga ggaatctcaga ccagagctcc aggaagttct gctggttggtc 180

tgataccaag agtaccttca gattctggaa aggattttca cgggggttgcc agtccagaat 240

tctttgctcc tcaaggctgt acccag 266

<210> 886

<211> 402

<212> DNA

<213> Homo sapien

<400> 886

cgcgtggttt ccgattgttt gatagtattt actggagaga tcatagaaac gactgtgaac 60

cgatgtcaca ccaggaaggt tgttgaqcat ttcttcaaca tcttcaattg tttcctttgt 120

aacctgtaag tccccgatgt ttaatttttag agctccaatt gctgttttac acaggatcac 180

tgcctcatca ttactttttca ccttctcacg agtcttttcc agaaaagtaa gagccacatt 240

aggatcagtc atctgtctaa ctacatgaag aatgatttcc acgagggaca aagggttcac 300

cctgtgttca aattcactga taaagttttc ataaagctta atgagaccat ctccttgggc 360

aaagcacgga tcctgcacaa aatcaagcac ctgaagtgtc ag 402

<210> 887

<211> 342

<212> DNA

<213> Homo sapien

<400> 887

ccaaagcgaq aqcattggca gtgaattgca gacactcttc cttgggtcatg ccttcccggc 60

aggtagcatc aacataacca tagatgtagg agctcccgga gcctccaatg gcaaaggact 120

gccttaccat catacccccc ataggcactg agtacacctg ccctccttct tgaggggtccc 180

agcctgcgat gatgattccc gccatcaggt cttcccggta tcggtaacac atctccttaa 240

agaagctggc tgcctgtgtgg accagtggag gctcattcag ttcaatgctg tggaaaccga 300

gctggtaggt gacagcatca gctactgcct gggatatcagc ag 342

<210> 888

<211> 228

<212> DNA

<213> Homo sapien

<400> 888

cgcgtcggcc aaggctgctg ctgttgctcc tccaaagaag gttggcttca aggccgtgtc 60

caqqqaccca cgaqcagagg cactggggggg caagggatct ccaaggggggc aagggatccc 120

taaagggggt agctcacagg tgaggggggt tagggcccct ctagggagcg cctgaggcca 180
tacattcaag agtgtccctg gtgaggccca ggaagagcc aggactgg 228

<210> 889
<211> 378
<212> DNA
<213> Homo sapien

<400> 889
ttggcttttc tcccccttc atcctcctct cccctttcct cactgaaggc tgtgagttgc 60
tttcaatgtg acaacactat gatgtcattt ggaaggattt gccaggacag actgattctg 120
agtctctgggt gccgtatgtg tatgcggcag tgttgtcagg cgatcttgtt tgaagctcta 180
tgttgccata attaccatca agtacacact gttggcaaaa ggctaacacc tgactttagg 240
aaatgctgat ttgagaacaa aaggaaagggt cttttttcac tgcttaaagt ggggtcactt 300
tgataccttt gcggtcatgt ctgtgtctga tgagtgtaga atctctggat gtgcactgtc 360
agtcattgtgt ccaccagg 378

<210> 890
<211> 215
<212> DNA
<213> Homo sapien

<400> 890
ccatttttga gtgtgtccat tgggtagcaa tgtggaaacc accagggcct ttgtggagaa 60
aatggagggg gttgagggag tcccaggagg ggcttatattg agggcctttg ccacttgctc 120
ataggcgagc tcgatctcct catcatctgg acaggtggaa gcgaattctt cccgggcgta 180
ggcattgctc aagtaccgat gcactccccg gaagg 215

<210> 891
<211> 412
<212> DNA
<213> Homo sapien

<400> 891
ctgggtcaagt tcaacagagc cttggctgac cattctatgg ctcaggcacc tcggctcatt 60
gatggcattg ttctttaccaa atttgatacc attgatgaca aggtgggagc tgctatttct 120
atgacgtaca tcacaagcaa acccatcgtc tttgtgggca ccggccagac ctactgtgac 180
ctacgcagcc tcaatgccaa ggctgtgggt gctgccttca tgaaggctta acgtggctct 240
tgcccaatac caaatcgccg ctttccccac aagcccttct tcctgtatca agaattgtgt 300
ttagagtatg tgagcaacct gtcttcagtg tagtacaaag gcagagttag ggggcttgtg 360
gtcctttcca accccactcc ccgttcagca cagccgccat ctgcaaggaa gg 412

<210> 892
<211> 472
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(472)
<223> n = A,T,C or G

<400> 892
tttttttttt tttttttttt ttaattacta ctttttattc taatgtgaac catggccctg 60

```

aaagctgata acaagcttgg ctgancagag ggaactaggg gtcggcagaa aggattatgg 120
gtggaaaaca ttggctcttc cttggggagt gatgctgggg aaaggggaana nagtggctca 180
ncctgcaggt aaataggcta naaaagccaa ggccaaaggc tggaggggag aggacagtca 240
gcatgtccag cctgggggtct ggggtgtagg ttatcccttc tccctgtgcc tccccatctc 300
gtccatgagc ctaggctcttg gagccttgtg ttggaggctg ctgtgatgtc aggaacgggg 360
atctgtctag cttttggcca cttcctggga cctcacgccc ctgttgacag atggagattg 420
ggcagcaggg ccttgctgcg ttgttatctg ctgttccgac ttggtttgtc tt 472

```

<210> 893

<211> 477

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (477)

<223> n = A,T,C or G

<400> 893

```

caaagattca ctttatttat tcattctcct ccaacattag cataattaaa gccaaaggagg 60
aggagggggg tgaggtgaaa gatgagctgg aggaccgcaa taggggtagg tccccgtgg 120
aaaaagggtc agaggccaaa ggatgggagg gggtcaggct ggaactgagg agcagggtgg 180
ggcacttctc cctctaacac tctccctgt tgaagctctt tgtgacgggc gagctcaggc 240
cctgatgggt gacttcgcag gcgtagactt tgtgtttctc gtagtctgct ttgctcagcg 300
tcagggtgct gctgaggctg taggtgctgt ccttgctgtc ctgctctgtg acactctcct 360
gggagttacc cgattggagg gcgttatcca ccttccactg tactttggcc tctctgggat 420
agaagttatt cagcangcac acaacanang cagtttccag atttcaactg ctcatca 477

```

<210> 894

<211> 289

<212> DNA

<213> Homo sapien

<400> 894

```

ctgtcttatg gctatgatga gaaatcaacc ggaggaattt ccgtgcctgg ccccatgggt 60
ccctctgggt ctcgtggtct ccctggcccc cctggtgcac ctggtcccca aggcttccaa 120
ggccccctg gtgagcctgg cgagcctgga gcttcaggct ccatgggtcc ccgagggtcc 180
ccaggtcccc ctggaaaagaa tggagatgat ggggaagctg gaaaacctgg togtcctggt 240
gagcgtgggc ctctggggcc tcagagtgtc cgaggattgc ccggaacag 289

```

<210> 895

<211> 179

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (179)

<223> n = A,T,C or G

<400> 895

```

ctggatgggt ccanacaaag tggaatccct ggaaccttta actgagcagt gaaggtcagt 60
gcctcagagc ctgagagatg aacaggacca gagagagagg tgggcaggca ggcacaagg 120
tatgtcttcc tcagactcgg aaccctgtct ttctccacca tccagacgtt cagctacag 179

```

<210> 896
 <211> 557
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1) ... (557)
 <223> n = A,T,C or G

<400> 896
 ccactcactg ctgggaccca ggcacctccc ttctccatcc tctctggatt gtcagtaatg 60
 tcctggaaca gaagcctgtg ggatggcctt gggcacggag aagccctggg gtcagtgtcg 120
 tgcacggatg gcggcagtgt tgaacccagg aggctgaacc cggccccacca cggaagatga 180
 gtgcatggca accgcctgcc ttcacgtcgc tccacttggg aacccccagg tctgggctgt 240
 tctaggtatt gcttcacgtg ccccagcaag cccttaacaa gagggcctgg ttcctgaag 300
 aaccaatccc aggaaggggc cttgatccct ccgccttgct gagagtgaac cctcgtctct 360
 cctcacnctc catttcattt ctgggaattg gggcttagtt tcgaaccttt ggcaaggctg 420
 ttcttactaa tgcccaagcc cctttacccc tctccctata ggttacacag gggagaccag 480
 ggccctcgga gaagactgct gccacacttc cgaatcattc tgcttgccaa ataggtcac 540
 ttcaccagtt gactgac 557

<210> 897
 <211> 495
 <212> DNA
 <213> Homo sapien

<400> 897
 ctggaatctc ctttgcaatc ccatctgata agattaaaaa gttcctcacg gagtcccatg 60
 accgacaggc caaaggaaga gccatcacca agaagaagta tattggatat cgaatgatgt 120
 cactcacgtc cagcaaagcc aaagagctga aggaccggca ccgggacttc ccagacgtga 180
 tctcaggagc gtatataatt gaagtaattc ctgatacccc agcagaagct ggtggtctca 240
 aggaaaacga cgtcataatc agcatcaatg gacagtccgt ggtctccgcc aatgatgtca 300
 ggcacgtcat taaaagggaa agcacctga acatggtggt ccgcaggggt aatgaagata 360
 tcatgatcac agtgattccc gaagaaattg acccataggc agaggcatga gctggacttc 420
 atgtttccct caaagactct cccgtggatg acggatgagg actctgggct gctggaatag 480
 gacactcaag acttt 495

<210> 898
 <211> 406
 <212> DNA
 <213> Homo sapien

<400> 898
 ccacgactgc atgcccgcgc ccgccagggtg atacctccgc cggtgaccca ggggctctgc 60
 gacacaggga gtctgcatgt ctaagtgcta gacatgctca gctttgtgga tacgcggact 120
 ttgttgctgc ttgcagtaac cttatgccta gcaacatgcc aatctttaca agaggaaacc 180
 gtaagaaagg gcccagccgg agatagagga ccacgtggag aaaggggtcc accaggcccc 240
 ccaggcagag atggtgaaga tgggccaca ggccctcctg gtccacctgg tcctcctggc 300
 cccctgggtc tcggtgggaa ctttgctgct cagtatgacg gaaaaggagt tggacttggc 360
 cccggaccaa tgggcttaat gggacctaga ggccacctg gtgcag 406

<210> 899

```
<210> 903
<211> 228
```

<212> DNA

<213> Homo sapien

<400> 903

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctggagactc | tgggccagga | gaagctgaag | ctggaggcgg | agcttggcaa | catgcagggg | 60 |
| ctggaggagg | acttcaagaa | caagtatgag | gatgagatca | ataagcgtac | agagatggag | 120 |
| aacgaatttg | tcctcatcaa | gaaggatgtg | gatgaagctt | acatgaacaa | ggtagagctg | 180 |
| gagtctcgcc | tgggaagggt | gaccgacgag | atcaacttcc | tcaggcag | | 228 |

<210> 904

<211> 388

<212> DNA

<213> Homo sapien

<400> 904

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaagcgctc | agatcggcaa | ggggcaccag | tcttgatctg | cccagtgcac | agccccacaa | 60 |
| ccaggtcagc | gatgaaggta | tcttcagtct | ccccgaacg | atgaggcacc | atgacgcccc | 120 |
| aaccattggc | ctgggccagc | ttgcacgcct | gaagagactc | ggtcacggag | ccaatctggt | 180 |
| tgactttgag | caggaggcag | ttgcaggact | tctcgttcac | ggccttggcg | atcctctttg | 240 |
| ggttggtcac | tgtgagatca | tccccacta | cctggattcc | tgcactggct | gtgaacttct | 300 |
| gccaaagctc | ccagtcaccc | tggtaaagg | gatcttcgat | agacaccact | gggtagtcct | 360 |
| tgatgaagga | cttgtacag | tcagccag | | | | 388 |

<210> 905

<211> 272

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(272)

<223> n = A,T,C or G

<400> 905

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccggagccca | cggnggtcat | ggctgccaga | gcgctctgca | tgctggggct | ggtcctggcc | 60 |
| ttgtgtctct | ccagctctgc | tgaggagtac | gtgggcctgt | ctgcaaacca | gtgtgccgtg | 120 |
| ccagccaagg | acagggtgga | ctgcggctac | ccccatgtca | cccccaagga | gtgcaacaac | 180 |
| cggggctgct | gctttgactc | caggatccct | ggagtgcctt | ggtgtttcaa | gccccctgcag | 240 |
| gaagcagaat | gcaccttctg | aggcacctcc | ag | | | 272 |

<210> 906

<211> 525

<212> DNA

<213> Homo sapien

<400> 906

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtgcaccc | gagtgtcctt | ccccccctaa | gctggcacat | aggagcaaaa | gttcactaac | 60 |
| cctgcagtgg | aaggcaccaa | ttgacaacgg | ttcaaaaatc | accaactacc | ttttagagtg | 120 |
| ggatgagggg | aaagaaatag | tggtttcaga | cagtgtctct | tcgggagcca | gaagcactgc | 180 |
| aagttgacaa | agctttgtcc | ggcaatgggg | tacacattca | ggctggccgc | tcgaaacgac | 240 |
| attggtacca | gtggttatag | ccaagagggt | gtgtgctaca | cattaggaaa | tatccctcag | 300 |
| atgccttctg | caccaaggct | ggttcgagct | ggcatcacat | gggtcacgtt | gcagtggagt | 360 |
| aagccagaag | gctgttcacc | cgaggaagtg | atcacctaca | ccttggaat | tcaggaggat | 420 |
| gaaaatgata | accttttcca | cccaaaatac | actggagag | atttaacctg | tactgtgaaa | 480 |

aatctcaaaa gaagcacaca gtataaattc aggctgactg cttct

525

<210> 907

<211> 365

<212> DNA

<213> Homo sapien

<400> 907

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| gtaaatttta | agtctttcag | ttttatagat | acggaaaaca | agggtgactc | tttaccacag | 60 |
| gatgaataaa | gaactaagta | atatgggaaa | tgcagcaatt | tctggactag | ctgagccgat | 120 |
| tccttctctgt | gagcacactg | taagctttca | agttctctgg | gcaggaatta | cagcacctgt | 180 |
| ccccgtcaat | ggccctgctg | tgtgatgctc | atcgcttccc | tctgtgctgg | agcagtcccc | 240 |
| caggtgtcca | tctcctatct | ttttgttcca | atcttctgtg | agttccagct | agcaggcttt | 300 |
| acatctgggg | aaaggaaaac | caggggtttt | agctctgttc | tctgctccca | tccttcgctc | 360 |
| accag | | | | | | 365 |

<210> 908

<211> 608

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(608)

<223> n = A,T,C or G

<400> 908

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| cggaggtgcc | tcagccatgg | catggatccc | tctcttctc | ggcgctcttg | cttactgcac | 60 |
| aggacgtgcg | gcctcctttg | aggtgaccca | gccaccttca | atgtccgtgt | ccccaggaca | 120 |
| gacagccaag | atcacctgca | ctggagatag | gttgggggat | gaatatgttt | gctgggtatca | 180 |
| acagaagcca | ggccagtccc | ctgtattgat | aatataatttg | gataacaagc | ggccctcggg | 240 |
| gatccctgac | cgattctctg | cctacgcctc | tgggaacaca | gccactctga | tcatcagcgg | 300 |
| ggcccaagtt | atggatgagg | cttattatta | ctgtcaggcg | tgggacggca | gaactgtggt | 360 |
| gttcggcgaa | gggaccaacc | tgaccgtcct | aggtcagccc | aaggetgccc | cctcgggtcac | 420 |
| tctgttcccg | ccctcctctg | aggagcttca | agccaacaag | gccacactgg | tgtgtctcat | 480 |
| aagtgacttc | taccggggag | ccgtgacagt | ggcctggaag | gcagatagca | gccccgtcaa | 540 |
| ggcgggagtg | gagaccacca | caccctccaa | acaaagcaac | aacaagtacg | cggncagcag | 600 |
| ctatctga | | | | | | 608 |

<210> 909

<211> 513

<212> DNA

<213> Homo sapien

<400> 909

| | | | | | | |
|-------------|------------|-------------|------------|------------|-------------|-----|
| ctgggtctcaa | actcctcacc | tcaactgatc | cgcccacctt | ggcctcccaa | agtgtctggga | 60 |
| ttataggtgt | gagccaccgt | gccc aaagtt | aagtattttt | gatcaagtgt | tttgtctttt | 120 |
| gtgcaaggca | tttgtggctc | tgtcatagca | gaggaaaaca | aaacatgcct | atcaaatgaa | 180 |
| tcaagtccga | cctcttctca | tattgagcaa | ctagaggctc | aggaacattt | cccctacctg | 240 |
| tcattctcat | ctggcatacc | aggtgtacat | actccttctt | attctcctct | gttaccaaga | 300 |
| tgttggtcccc | attgggtttg | aggtcacgaa | ctccacaaac | tccaaactct | tggacctcag | 360 |
| tgtgtgaagg | gaggtcatag | cctagtgtgg | agacatcatt | ttccagcaga | taaaccagac | 420 |
| cttggtagaa | gtggtaatct | tcactctcca | tatctgtata | tctgactgac | ttgcccaaga | 480 |
| tgtgtttgta | aaaggatcga | gtaaagtagc | act | | | 513 |

| | | | | | | | | |
|------------|------------|------------|------------|-------------|------------|--|--|-----|
| <400> | 913 | | | | | | | |
| cctggacacc | ataaggctgg | tgggctttca | gaattgtgtt | agggggggcag | gagtggcagg | | | 60 |
| ttcctgaatc | tcggtcaata | tagtaaccag | caggacaaga | ggtgcaggag | gagcccat | | | 120 |
| cagaggcttc | tagggcacag | ggacggcagt | aggaggccac | gccattcata | acattggtga | | | 180 |
| cattgatgga | gtagatcttg | gcaacgtcat | tgggtgactt | cctgcttgcc | tcatgaaaag | | | 240 |
| tggtcctctg | gaaggcccag | gtgaggctcg | tggtagtggt | ctcctcaatg | atgtaggtat | | | 300 |
| aggactgttt | gcctttggaa | cctttccacg | tctccacagg | agtgttggtc | ctagaattca | | | 360 |
| caccaccat | gaagtagagc | tcacagttca | cagaacagag | ggtctcaaag | acaaatgtga | | | 420 |
| ttctgg | | | | | | | | 426 |

| <400> 917 | | | | | | | |
|------------|------------|------------|------------|-------------|------------|--|-----|
| ggcacagcga | gggcagcatc | tggaggagct | ctgcagcctc | cacacctacc | acgacctccc | | 60 |
| agggctgagc | tcaggaaaaa | ccagccactg | ctttacagga | caggggggttg | aagctgagcc | | 120 |
| cgcctcaca | cccaccccca | tgcactcaa | gattggattt | tacagctact | tgcaattcaa | | 180 |
| aattcagaag | aataaaaaat | gggaacatac | agaactctaa | aagatagaca | tcagaaattg | | 240 |
| ttaagttaa | ctttttcaaa | aaatcagcaa | ttccccagcg | tagtcaaggg | tggacactgc | | 300 |

```

acgctctggc atgatgggat ggcgaccggg caagctttct tctcgagat gctctgctgc 360
ttgagagcta ttgctttgtt aagatataaa aaggggtttc tttttgtctt tctgtaagg 420
ggtcttccag cttttgattg aaagtcctag ggtgattcta tttctgctgt gatttatctg 480
ctgaaagctc ag 492

```

<210> 918

<211> 557

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(557)

<223> n = A,T,C or G

<400> 918

```

ctgctcctgg gtaggcgtgc gggccatata gtaggggtag gatactagcc gctcgccgcc 60
gttcagattt gctcccagca cgaagggtt cttctccatc caggcaatga tggcccggac 120
ctccgtggat accgtggcat ctggcgaaag gtagcggtca gggatgggca agttattgtt 180
ggggaccggg taggggacct atttctctc ctcagctccc cagagcacag agttgagatc 240
cgggaaatct tcaaagatgt caaagccctc ctcagtccac agtcccagcg occagttccc 300
aaactctgag cccatctgag ctgccacctc gtagccatca gggttcagtg agggcaccag 360
gtggatgcgt gtgtcctgca ccaggctgag cacacgtggg ttcccatcgc ggtactctcg 420
gcacaggtac tgcattgagc gcagcaacag ctctcgcccc agcacctcgt tgccatggat 480
cccagcagtg tagcggaact cgggctcccc cagttcatgc tcccanggt tgtctgagat 540
ctccatggca tagatct 557

```

<210> 919

<211> 407

<212> DNA

<213> Homo sapien

<400> 919

```

ccttatgact acaacggccc acgagaaaaa tatggaatcg ttgattacat gatcgagcag 60
tccgggcctc cctccaagga gattctgacc ctgaagcagg tccaggagtt cctgaaggat 120
ggagacgatg tcatcatcat cggggctctt aagggggaga gtgaccagc ctaccagcaa 180
taccaggatg ccgctaacaa cctgagagaa gattacaaat ttcaccacac ttctatcaca 240
gaaatagcaa agttcttgaa agtctcccag gggcagttgg ttgtaatgca gcctgagaga 300
ttccagtcta agtatgagcc ccggagccac atgatggagc tccagggtc caccagggac 360
tcggccatca aggacttcgt gctgaagtac gcctgcccc tgggttgg 407

```

<210> 920

<211> 340

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(340)

<223> n = A,T,C or G

<400> 920

```

cctcttgggc agcnnagggc cctgcctctg tttcatgatg catgggtcat ttgtcttggg 60
tgtcctatcc catatggaga agaaaggggc tctaagttct ggctcttctt tctttggggt 120

```

```
<210> 921
<211> 571
<212> DNA
<213> Homo sapien
```

```
<210> 922
<211> 262
<212> DNA
<213> Homo sapien
```

| | | | | | | | | |
|------------|------------|------------|------------|------------|------------|--|--|-----|
| <400> | 922 | | | | | | | |
| gccaanaca | tncaggtcac | agcagattcg | ggcacgtgtg | gaagaaggtt | ggatgatgtc | | | 60 |
| atccacaaac | cctgcgactg | ctgcagggaa | agggttggca | aacttctcga | tgtactctgc | | | 120 |
| ctgancagct | tccacattct | cattgccctt | gaagatgatc | tccacagcgc | cctttgctcc | | | 180 |
| catgactgca | atctctgnng | tgggccangc | atanttggta | tcaccacaaa | ngtgcttaga | | | 240 |
| qctcatgaca | tcntaggcac | ct | | | | | | 262 |

| | | | | | | | | | |
|------------|------------|------------|------------|-------------|------------|--|--|--|-----|
| <400> | 923 | | | | | | | | |
| ccactgggac | tttggcttcc | tgatgccgat | tgtggatttc | tgctgcaaaag | acagtgatgt | | | | 60 |
| tgagccaggc | tgtttcctct | ctatccagag | gttttgtagt | tttaataaaa | ccatcctctg | | | | 120 |
| gattaatagt | gaaaaatctg | tcgaggtcag | tgtgacgadc | gatggaatac | cttatcgggc | | | | 180 |
| tgttggcagc | atcagggtct | ttggcatgca | ctctcccaac | cacggtgccca | gcag | | | | 234 |

```
<210> 924
<211> 152
<212> DNA
<213> Homo sapien
```

<400> 924
ccaggattga caggccatcc attcacagcc aggagatgct gggccagttc ctccaagagg 60
tctccgtcat ggcagtgatg aaaacctaac aggggtggccc cctgtgccag ctcaggtgac 120
tggagcccga gggcctgaca ggttcccagc ag 152

<210> 925
<211> 400
<212> DNA
<213> Homo sapien

<400> 925
caatatcatg ccaaggaccc aaacaacctc ttcattggtgc gcttggcaca gggcctgaca 60
catttaggga agggcaccct taccctctgc ccctaccaca gcgaccggca gcttatgagc 120
caggtggccg tggctggact gctcactgtg cttgtctctt tcctggatgt tcgaaacatt 180
attctaggca aatcacacta tgtattgtat gggctgggtgg ctgccatgca gcccgaatg 240
ctggttacgt ttgatgagga gctgcggcca ttgccagtgt ctgtccgtgt gggccaggca 300
gtggatgtgg tgggccaggc tggcaagccg aagactatca cagggttcca gacgcataca 360
accccagtgt tgttggccca cggggaacgg gcagaattgg 400

<210> 926
<211> 521
<212> DNA
<213> Homo sapien

<400> 926
ccacgtccct attttagaaa tgagaggagt gactgcacac aggaaaaatg ccacttttag 60
caattcaaag tggaaaaact tcttttatat aaaaattatc ccaactccca ccccttggct 120
ctcagtgttg catctcccac agaggtaaag ttgtgccatt ttcccacggc tttaaacaaa 180
gcaaaacaaa accaccaatc ctaataacct ccctccctgc cccgtctcca cgctgtgcgg 240
agagggtctt agcccctcag tcggaettct ccttctcctt catgtgcaag aagacgatgc 300
tgaagatgaa gagccccagc atcatggaga aggcgctggc gtagtagggg taggcccagg 360
ggatgaagcg ctcatactgc gtgtgctgga gtggccgcac ggatacctga gtggaagagt 420
acaggtgtgt gtagcctagc cggttgaat ccactttaaa ctggaataca ccatacacgt 480
cgggcaactt gaactgaaca ctgtatttgc cacctttctt c 521

<210> 927
<211> 520
<212> DNA
<213> Homo sapien

<400> 927
ccaggctagt ctcgaaactc tgacctcagg tgatctgcct gcctcggcct cccaaagtgc 60
tgggattacc ggcgtgagcc accatgcctg gccttacatt ttttaaaatg agggaaacaaa 120
tgaataaatg accaccatgt taggggctgg ctctgaacag aattgtaaag tgggccaagc 180
ttgctctcaa ggtcacctta agcccacggt tgctgtgtcc tgccctctca gggtcatttc 240
ccagcctcca ggcacctgtt cacagaggct gcactctggc tcgcctccac ccctccatcc 300
taaggtgctc cgtgactta gaacaggaca gtcagggaga gaatgtgtct caggagggtg 360
gagtcagatg atcacggcct tcctggcatc tgaggggata cagcttcggg tagcaaagtg 420
tgattttccc tgagccccag gaaagcttgg ccttggctag aatacattga accctgaggg 480
ccagagagtc cctggggcaa gctctgagag ggaggacctc 520

<210> 928
<211> 492

<212> DNA
<213> Homo sapien

<400> 928
ctgagcttttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa 60
agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatatac ttaacaaagc 120
aatagctctc aagcagcaga gcatctcgag gaagaaagct tgcccggctc ccatcccatc 180
atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa 240
agcttaactt aacaatttct gatgtctatc ttttagagtt ctgtatgttc ccatttttta 300
ttcttctgaa ttttgaattg caagtagctg taaaatccaa tctctgagtg catgggggtg 360
gggtgtgaggc ggggctcagc ttcaaccccc tgctcctgtaa agcagtggtt ggtttttctt 420
gagcccagcc ctgggaggtc gtggttaggtg tggaggctgc agagctcctc cagatgctgc 480
cctcgctgtg cc 492

<210> 929
<211> 209
<212> DNA
<213> Homo sapien

<400> 929
ttttttcacc atctaacaaa ggcactttat tgcattacca ttcacaatta acagtcaaga 60
acaaataata ataacaaata aaataacttt taagaggaca aggcattaga aataaaaaag 120
gacactaata acatttgtaa aagcttgtagc tggatgtggt tgccccatt tgtgtgtgtg 180
gttgtgtgtg tgtggttgtg tgttggtgg 209

<210> 930
<211> 617
<212> DNA
<213> Homo sapien

<400> 930
cgcgtccttt aacaagcccc gttctcaaaa ggctgggggt atttatataa gaacttattc 60
caaagtgact ctaagatcca tggtcccaag atctagtacg ggctattcat gggtctgagg 120
catgtccagc atgcaggcaa acttatctgt tcaaattgag gtaaaacaga caaaaaacac 180
ttaatattaa cagaagctac ataattaaaa ctaaccttct gctgcttatt taagctaatt 240
atgtattctt accaaacaga gacctcaag tcaatcattt cttttgattt tagttaccac 300
ccccaaatta agcctcttct ttcaaagcca ttattagtta aaaaaaagtt ttaaaatgaa 360
gaaaaatatt ttttcagaa cttgtatttt gtaattagtg tgatgcaatt tctttttatt 420
tttcaaactt agaaataact catgtatggt actatttggt atttttttca gataccaagg 480
aataccgaca ggattcataa ataggatttt ctgacactgg caggaaagtc tgctaacgtt 540
tacaaaatac caaagactct tctttcaagc ttcaaagatg gctgagaatt aacagttatg 600
attagttttt cagtaca 617

<210> 931
<211> 521
<212> DNA
<213> Homo sapien

<400> 931
ccaacaaaat tgggtgaacac atggaagaac atggcatcaa gtttataaga cagttcgtac 60
caattaaagt tgaacaaatt gaagcagga caccaggccg actcagagta gtagctcagt 120
ccaccaatag tgaggaaatc attgaaggag aatataatac ggtgatgctg gcaataggaa 180
gagatgcttg cacaagaaaa attggcttag aaaccgtagg ggtgaagata aatgaaaaga 240
ctggaaaaat acctgtcaca gatgaagaac agaccaatgt gccttacatc tatgccattg 300

```
<210> 932
<211> 197
<212> DNA
<213> Homo sapien
```

```
<210> 933
<211> 610
<212> DNA
<213> Homo sapien
```

```
<210> 934
<211> 384
<212> DNA
<213> Homo sapien
```

```
<210> 935
<211> 125
<212> DNA
<213> Homo sapien
```

<220>

$\langle 220 \rangle$

<221> misc_feature
 <222> (1)...(192)
 <223> n = A,T,C or G

<400> 938
 tttttttttt tttttttttt ttttttttngg aaaaagccca aaaggcactt tattggaggt 60
 ctntgcctcc attcacagga aaaaggagct gggagcccca tcctaagggt cccagcatca 120
 gccactgga gggcctggaa cagtccanca ctntgtggga aaggagtggg gaggggaatg 180
 ttttaaaaaa aa 192

<210> 939
 <211> 337
 <212> DNA
 <213> Homo sapien

<400> 939
 ccaaaatatt ggaacacaca gaaccaaacc aggtgtgttc tacacctgca tgagtgaagg 60
 atttccacgt agacacctag gaagagcccg catgccttag actcactcca gaggaaggat 120
 tgatttgcaa ccagaaaggg agctgaaaac caccgagctc catggctctt cattcaaaag 180
 ggaaaataat gattccacgt tgcttttttag agttcaaadc aacatctttc tggataaatc 240
 tattttttta caatcttttt attatttgta aaagatataa aaacaactoc catcagtagc 300
 aatacaagggt tatacatttt aaccagattt tctcagg 337

<210> 940
 <211> 362
 <212> DNA
 <213> Homo sapien

<400> 940
 cctgtccaaa cgtgcgcacc aggaccgagg ggagctccct cccaacacct gctaggaatt 60
 gccaaactttt aaatggatgg ggttttttat ggggtgaacc tctgttaata cttttgtaca 120
 ctctcactac agtttatatt tttataggct attttctcaa ggtgtttcta gattccacat 180
 atctatttta tataacaagt tattatgtta tgtgtgtgac tcccttgtgt gtatctgtgc 240
 cagcctcagc ctccgagttg cttttccctc tggccctgac tctcactgac tcaccgatgt 300
 ggtgtgcagg cccacttctt accccagata gctcggggcg ctgcctgtag tcatgccgac 360
 ag 362

<210> 941
 <211> 216
 <212> DNA
 <213> Homo sapien

<400> 941
 ctggacatct ttccagcccg ggatacctac catcctatga gcgagtaccc cacctaccac 60
 acccatgggc gctatgtgcc ccctagcagt accgatcgta gccctatga gaagggtttct 120
 gcaggtaatg gtggcagcag cctctcttac acaaaccag cagtggcagc cactttctgcc 180
 aacttgtagg ggcattgtgc ccgctgagct gattgg 216

<210> 942
 <211> 324
 <212> DNA
 <213> Homo sapien

<400> 942

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgattggct | tcaggccccc | tacctctata | aactctacca | gcattactac | ttcctggaag | 60 |
| gtcaaattgc | catectatat | gtctgtggcc | ttgcctctac | agtcctcttt | ggcctagtgg | 120 |
| cctcctccct | tgtggattgg | ctgggtcgca | agaattcttg | tgtcctcttc | cccctgactt | 180 |
| actcactatg | ctacttaacc | aaactctctc | aagactactt | tgtgctgcta | gtggggcgag | 240 |
| cacttggtgg | gctgtccaca | gcctgtctct | tctcagcctt | cgaggccagg | gagcctcaaa | 300 |
| tcttcagtct | ctcagagacc | acag | | | | 324 |

<210> 943

<211> 597

<212> DNA

<213> Homo sapien

<400> 943

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| ctgacaaaat | tcctgggtta | ctaggtgtct | ttcagaagct | gattgcatcc | aaagcaaattg | 60 |
| accaccaagg | tttttatctt | ctaaacagta | taatagagca | catgcctcct | gaatcagttg | 120 |
| accaatatag | gaaacaaatc | ttcattctgc | tattccagag | acttcagaat | tccaaaacaa | 180 |
| ccaagtttat | caagagtttt | ttagtcttta | ttaatttgta | ttgcataaaa | tatggggcac | 240 |
| tagcactaca | agaaatat | gatggtatac | aacaaaaaat | gtttggaatg | gttttggaat | 300 |
| aaattattat | tcctgaaatt | cagaaggtat | ctggaaatgt | agagaaaaag | atctgtgcgg | 360 |
| ttggcataac | caaattacta | acagaatgtc | ccccaatgat | ggacactgag | tataccaaac | 420 |
| tgtggactcc | attattacag | tctttgattg | gtctttttga | gttaccgcga | gatgatacca | 480 |
| ttcctgatga | ggaacatttt | attgacatag | aagatacacc | aggatatcag | actgccttct | 540 |
| cacagttggc | atttgctggg | aaaaaaagag | catgatcctg | taggtcaaatt | ggtgaat | 597 |

<210> 944

<211> 359

<212> DNA

<213> Homo sapien

<400> 944

| | | | | | | |
|------------|------------|------------|-------------|-------------|-------------|-----|
| ctggaagagg | aaaaggagat | actgcagaaa | gaactctctc | aacttcaagc | tgacacaggag | 60 |
| aagcagaaaa | caggtactgt | tatggatacc | aaggctcgatg | aattaacaac | tgagatcaaa | 120 |
| gaactgaaag | aaactcttga | agaaaaaacc | aaggaggcgag | atgaataactt | ggataagtac | 180 |
| tgttccttgc | ttataagcca | tgaaaagtta | gagaaaagcta | aagagatggt | agagacacaa | 240 |
| gtggcccac | tgtgttcaca | gcaatctaaa | caagattccc | gagggctctcc | tttgctaggt | 300 |
| ccagttgttc | caggaccatc | tccaatccct | tctgttactg | aaaagaggtt | atcatctgg | 359 |

<210> 945

<211> 367

<212> DNA

<213> Homo sapien

<400> 945

| | | | | | | |
|------------|------------|-------------|-------------|------------|-------------|-----|
| caggatctga | agtttggggg | cgagcaggat | gttgatatgg | tgtttgcgtc | attcatccgc | 60 |
| aaggcatctg | atgtccatga | agtttaggaag | gtcctggggag | agaaggga | gaacatcaag | 120 |
| attatcagca | aaatcgggaa | tcatgagggg | gttcggaggt | ttgatgaaat | cctggaggcc | 180 |
| agtgatggga | tcatggtggc | tcgtggtgat | ctaggcattg | agattcctgc | agagaagggtc | 240 |
| ttccttgctc | agaagatgat | gattggacgg | tgcaaccgag | ctgggaagcc | tgatcatctgt | 300 |
| gctactcaga | tgctggagag | catgatcaag | aagccccgcc | ccactcgggc | tgaaggcagt | 360 |
| gatgtgg | | | | | | 367 |

<210> 946

<211> 335

<212> DNA

<213> Homo sapien

<400> 946

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ccacagaggt | ggtattacaa | aatatacaaa | gtgggtttctt | tctttacatt | tcatagaaga | 60 |
| agcctgcctc | atttccaaat | gagagcacta | gaagcacaaa | tcatgcagac | catttactat | 120 |
| ataacttatg | aaaaatgctg | tacagggctg | tgactataga | tatagagtat | ttggctctgt | 180 |
| ttgggaattg | atatctacaa | gggggagggg | caggggagga | ctgtccgata | tcctgacttg | 240 |
| ctgggatggt | ggagaagctg | ggatggggga | ggccccaatc | ttgctgcacg | gctacacca | 300 |
| ctcctccttt | cctagacaag | gctggagcgc | actgg | | | 335 |

<210> 947

<211> 384

<212> DNA

<213> Homo sapien

<400> 947

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| cctcttgag | cacatccttt | actgcattgt | ggacagcgag | tgtaagtcaa | gggatgtgct | 60 |
| ccagagttac | tttgacctcc | tgggggagct | gatgaagttc | aacgttgatg | cattcaagag | 120 |
| attcaataaa | tatatcaaca | ccgatgcaaa | gttccaggta | ttcctgaagc | agatcaacag | 180 |
| ctccctgggtg | gactccaaca | tgctggtgcg | ctgtgtcact | ctgtccctgg | accgatttga | 240 |
| aaaccaggtg | gatatgaaag | ttgccgaggt | actgtctgaa | tgccgcctgc | tcgcctacat | 300 |
| atcccaggtg | cccacgcaga | tgtccttcct | cttccgcctc | atcaacatca | tccacgtgca | 360 |
| gacgctgacc | caggagaacg | tcag | | | | 384 |

<210> 948

<211> 173

<212> DNA

<213> Homo sapien

<400> 948

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ctgtggaggg | gacactgtct | ttgaggcatc | actggttcca | caaagggtag | gggaaggctc | 60 |
| tgaggggacca | ccccatgcc | tcattaatca | accagaagct | tggcctggag | cagcagcggg | 120 |
| gattccagta | gctgtgggca | tacaggatgc | tagggcggcc | acaaccagg | cag | 173 |

<210> 949

<211> 211

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (211)

<223> n = A,T,C or G

<400> 949

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ccatccacgt | tgnaaacag | aataaaatgg | aaattcacct | tgatcatctac | ccgacattgg | 60 |
| ccttctgtg | ccacggcatc | atgggctgcc | tgtatggcct | cattcttttc | aaagcatttt | 120 |
| gctctgtctt | caggggacat | tttctctgtt | tcagaaagaa | actgtttcag | aactgatcca | 180 |
| tcctcaaate | ccagtttgtc | ttgattattg | g | | | 211 |

<210> 950

<211> 382

<212> DNA

<213> Homo sapien

<400> 950
 cctcatcggtg agtcaggacg tgggtgaaagc tgcagtggct gctgtgctct ctccagaaga 60
 attcatgggtc ctgttggact ctgtgcttcc tgagagtgcc catcggctga agtcaagcat 120
 cgggctgac aatgaaaagg ctgcagataa gctgggatct acccagatcg tgaagatcct 180
 aactcaggac actcccgagt tttttataga ccaaggccat gccaaagggtg cccaactgat 240
 cgtgctggaa gtgtttccct ccagtgaagc cctccgccct ttgttcaccc tgggcatcga 300
 agccagctcg gaagctcagt tttacaccaa aggtgaccaa cttatactca acttgaataa 360
 catcagctct gatcggtacc ag 382

<210> 951
 <211> 473
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1) ... (473)
 <223> n = A,T,C or G

<400> 951
 cctctctgcc aggcaaagga gggagctgcg gctctttgac attaaaccag agcagcagag 60
 atacagcctt ttcctccctc tccatgaact ctggaaacag tacatcaggg acctgtgcag 120
 tgggctcaag ccagacacgc agccacagat gattcaggcc aagctcttaa aggcagatct 180
 tcacggggct attatttcag tgacaaaatc caaatgcccc tcttatgtgg gtattacagg 240
 aatcctttcta caggaaacaa agcacatttt caaaattatc accaaagaag accgcctgaa 300
 agttatcccc aagctaaact gcgtgttcac tgtggaaacc gatggcttta tttcctacat 360
 ttacgggagc aaattccagc ttcggtcaag tgaacggtct gcgaagaagt tcaaagcgaa 420
 nggaacgatt gacctgtgaa ttctttgccg tctaangcag ttgtttatga cag 473

<210> 952
 <211> 312
 <212> DNA
 <213> Homo sapien

<400> 952
 ctgatgggtc tcatagtcct ctgggatggg gtcattgcag cggtaacgca gggttgccca 60
 gatgatgttc tcctgggaga agcagaagac ccccaagcgg ccaccccgca tggttgtgtc 120
 caagaccacg ttgctgtcgg ccaccagctc agggccctca tagaatcgca ccctgatgta 180
 gccactttgg ggccggtgct gcaggaacca acgataggac ttcttgtcct tccaaccac 240
 gtttcgctgg tccttcaca gcagccgcac ctgagactct gtgtctcctg tatgccacag 300
 agcgttccgc ag 312

<210> 953
 <211> 397
 <212> DNA
 <213> Homo sapien

<400> 953
 cgcgtccact gccgaccctc ttggtttctg aaaccaacct ttcttcctgc tctcctcttt 60
 aagagcaaac cccaacatgt ataaggctac agcaagtggg agccaggaaa agctgtggga 120
 cccctcatct gagtcacatc catatggcat ggagaaagaa aacctctctg ccagaaggaa 180
 ctgaactctg gaagtcctaa ggaaggctac catgatcagc agataggaaa gcattgccaa 240
 gggctgtccc tcaagagctt agttttctta gggagaccag aaagacatca gatcctgact 300

```
<210> 954
<211> 304
<212> DNA
<213> Homo sapien
```

```
<210> 955
<211> 156
<212> DNA
<213> Homo sapien
```

```
<210> 956
<211> 543
<212> DNA
<213> Homo sapien
```

```
<210> 957
<211> 528
<212> DNA
<213> Homo sapien
```

| <400> 957 | | | | | | | |
|------------|------------|------------|-------------|------------|-------------|--|-----|
| ctgtgatcaa | gatgtattaa | agaatatga | aagagcatct | gggttattct | agaagttctg | | 60 |
| tgatcaaaac | atattaaaaa | aaattaaagc | gcattctgggt | tattctagaa | gttctctgggc | | 120 |
| tttatacttg | gatatttaca | gaggaagttg | aacttcaagt | tctgccactc | ttcaaaatgg | | 180 |
| gtgacaggag | aggacgtgat | aggacagtta | aaaaaaaatt | gatagtcatt | ctctgatgga | | 240 |
| gtgaagcaag | ctttgtcaac | catcaacaaa | tatgaacttca | ttgggtcaca | gccttcgaga | | 300 |

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| gatccaacaa | gatttgagtt | ttaaatacag | aacatatttc | aaacagaacc | agcagagtgc | 360 |
| tgatgtatga | atggaattga | ttgctgaagg | cagagagtat | aaagaatctc | aagaaacttt | 420 |
| tagtgccatt | ttcattttaat | aagccattgg | tatagcaacc | taaaaacctt | ggctgtgatg | 480 |
| acaccaggat | gtgtttatgg | aattgctgca | ggagaacaca | attggcag | | 528 |

<210> 958

<211> 451

<212> DNA

<213> Homo sapien

<400> 958

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ctgtctgacc | atggggacct | tctgtctgaa | gaggagctgg | atgaatgaga | ctctgggaat | 60 |
| catctacaca | ggaccaaacc | caacaggcgc | cctggcaccg | gggaggcggg | tagttgtact | 120 |
| ctgcttgtag | agtccttgag | cccagtttac | agatctggag | agcaggaggc | caggacaagg | 180 |
| acaaaggctg | gaggatggag | taggacccag | gggctctgcc | atcctaggca | tcattcaagg | 240 |
| tcttttatga | agactttaca | gatgtcctct | gtaagtagca | tcgagagtgg | agttcagctc | 300 |
| ctttctctac | ttttttttgg | tctgatggca | catatttatt | gttctgtggt | ctaatacacag | 360 |
| tgtttctaaa | tgtaaaaagt | gcatatgttg | gtgtagctag | tcccgcgaca | ttgagtcct | 420 |
| ctgcatgaag | acactgggct | cctgcatcca | g | | | 451 |

<210> 959

<211> 158

<212> DNA

<213> Homo sapien

<400> 959

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ccagaccaag | gctgctggac | ctatgggaat | attcggtgtg | ctgtagagga | tgtgactgtc | 60 |
| ctgggtggact | acacagtacg | gaagttctgc | atccagcagg | tgggcgacat | gaccaacaga | 120 |
| aagccacagc | gcctcatcac | tcagttccac | tttaccag | | | 158 |

<210> 960

<211> 235

<212> DNA

<213> Homo sapien

<400> 960

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgagcaggg | aatccggccg | gaggaaggag | cagcttaccg | actgcgggtg | ttcaccacag | 60 |
| gccaggccct | aatatgcacc | cactagttta | gctcagactc | ctctctacat | atgaatggca | 120 |
| aaggcacttt | tgatatacac | tgtaaaatac | actgtatttt | agaatcggaa | tctattttct | 180 |
| aatgttcccc | tcaagggtcg | agtggcagga | aggttgagga | tgcaggactt | tgcag | 235 |

<210> 961

<211> 375

<212> DNA

<213> Homo sapien

<400> 961

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| cctggaaaga | aaagggatat | gtccagcgac | ttggagagag | accatcgccc | tcatgttagc | 60 |
| atgccccaga | atgccaacta | aactcctccc | tttcttccct | aatttccctt | cttgcatcct | 120 |
| tcctataact | tgatgcatgt | ggtttggttc | ctctctgggtg | gctctttggg | ctgggtattgg | 180 |
| tggctttcct | tgtggcagag | gatgtctcaa | acttcagatg | ggaggaaaga | gagcaggact | 240 |
| cacaggtttg | aagagaatca | cctgggaaaa | taccagaaaa | tgagggccgc | tttgagtccc | 300 |
| ccagagatgt | catcagagct | cctctgtcct | gcttctgaat | gtgctgatca | tttgaggaat | 360 |
| aaaattattt | ttccc | | | | | 375 |

<210> 962
 <211> 409
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(409)
 <223> n = A,T,C or G

<400> 962
 ctggggaggc ccnccgggcc tctcangtgg acaggtcacg gcattgggtg aagctggatg 60
 aagctggggc ctnnngctcct nctcatcaaa tacagatcac tnggaccctg tctcctcca 120
 tgggtgctggt ctctcgggcc ccaactgcccc tgetttctgt ttcttctctc acctcctcct 180
 cccccagctc catgtccagc tcgttgccctg cctctgaggg tgtgtagggt gagccactga 240
 tggaacggca gctaaagaag acgattcgct tgagccgctt gttgtagaag aagtagttga 300
 aggaccagag gctaccatcc tccccgaagg gatctgagtc caagtctggg ttatagctgt 360
 agatgtcaca ttcagccagg cagatctcct cgtccaccgc gttccacag 409

<210> 963
 <211> 163
 <212> DNA
 <213> Homo sapien

<400> 963
 gccatggcgt cctattttcga tgaacacgac tgcgagccgt cggaccctga gcaggagacg 60
 cgaaccaaca tgctgctgga gctcgcaagg tcaacttttca ataggatgga ctttgaagac 120
 ttgggggttg tagtagattg ggaccaccac ctgcctccac cag 163

<210> 964
 <211> 344
 <212> DNA
 <213> Homo sapien

<400> 964
 ccaactggctg agttattggc ctggcaggta tagagtccgc tgttcttctc agtgatgttg 60
 gagataaaga gctcttggtg gtgttgctgg atgttcccat caatcagcca agaatactgt 120
 gcagggtgggt tagaggctgc atggcaggag aggctgaggt tcaccctctg acggtaatag 180
 gtgtatgagg gggaaatggt ggggtcgtct gggccataga ggacattcag gatgactggg 240
 togtgtggt caacacttaa ttcgttctgg attccacact catagggtcc tacatcattc 300
 cttgtgacac tgagtagagt gagggtcctg ttgtcattgg acag 344

<210> 965
 <211> 461
 <212> DNA
 <213> Homo sapien

<400> 965
 ctgagctttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa 60
 agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatata ttaacaaagc 120
 aatagctctc aagcagcaga gcatctcgag gaaggaagct tgcccggctc ccatcccatc 180
 atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa 240
 agcttaactt aacaatttct gatgtctatc ttttagagtt ctgtatgttc ccatttttta 300

ttctttctgaa ttttgaattg caagtagctg taaaatccaa tctttgagtg catgggggtg 360
 ggtgtgaggc ggggctcagc ttcaaccccc tgtcctgtaa agcagtggct ggtttttcct 420
 gagcccagcc ctgggagggtc gtggtagggtg tggaggctgc a 461

<210> 966
 <211> 246
 <212> DNA
 <213> Homo sapien

<400> 966
 cctttcacag aactaccat tgagtgggtt gatgcagggt gcagccttca gtccccgagt 60
 actgggttct gataaaattc cacagaatcc agcatcactg ggctcagacg gcatccactg 120
 tagtaacta tttgtaaatg gggacatata tcccagcac cagtaggaca cattgatctt 180
 ccgaaggccg acccatgggg ttaagggtgag cttggacatg ctctgagatg actgcattat 240
 tcgcag 246

<210> 967
 <211> 244
 <212> DNA
 <213> Homo sapien

<400> 967
 ctggagcatt ggcagggaca gtcagaaagg agacaagtga aaacgggtcag atggacacag 60
 gcggaggaga aaagacagag ggagagagac catcggaac aatcagaggg gccgagacga 120
 tcagaaaagg gtcagcccga gacaggctga gccagagttt ctagaagcag tttccaattc 180
 aacggctcgc tttgagggcc aacgtgtcct aggccgaggc tgcagaagcg ctcacacact 240
 cagc 244

<210> 968
 <211> 436
 <212> DNA
 <213> Homo sapien

<400> 968
 ccaaagtctt taccctatctt aacccttctg atattttctga ctgctcactg ttcataattat 60
 aggggaccag atttgaata tagaattctc cataacatga atgaaattaa tgctgtccaa 120
 gccagcatgg tggcttcata ttaagtagta acagaagtct gaacaattgg ataaatttga 180
 cttccaagac agctaaactt ttcaactgca attttaaaaa ctacactaca ctgttatagt 240
 taatctgaca aaaatgtcct caaagagtac tttattttat ttaaagcatc tgtttaattc 300
 aacctttaat aattttgcaa agaagggtac gtgtgtattt taatatagcc tgacctgaat 360
 ttatatgttt ttagcttttag tatttaactt tttgtaacaa ataaaccttt tttaaaacaa 420
 gtttaaaaaa gaaaaa 436

<210> 969
 <211> 383
 <212> DNA
 <213> Homo sapien

<400> 969
 ctggctccct tgtctccagg gctttggagg atcagggtag ggagggctct gtctctaagc 60
 cagggtgtcag gatcagaatc atgggtagaa ggtgccattc agctcacagc cgcacccaga 120
 atcctttgca gccctccttc tttatttttt tccattgca ttctgggagt ccacatctgg 180
 ctttctcagc cactgttcat caccaggggt tttaggagga aggcttggct cctgtcttcc 240
 cagacccacc atgcctggag aggtcaggat ggaactacct cattcggcga attagcccca 300

aattgaacgc tgaatcgtgt cccatgagat caggcgccat ctgtaaagtc tcctctggaa 360
atgccaatcc atccttcccc cag 383

<210> 970
<211> 543
<212> DNA
<213> Homo sapien

<400> 970
ctgtagcttt tgtgggactt ccactgctca ggcgtcagge tcaggtagct gctggccgcg 60
tacttgttgt tgctttgttt ggaggggtgt gtggtctcca ctcccgctt gacggggctg 120
ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
agtgtggcct tgttggcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg 240
gcagccttgg gctgacctag gacggtcagc ctggctccctc cgcggaacac cgaagtgcta 300
ctgtttgtat atgagctgca gtaataatca gcctcgtcct cagcctggag cccagagatg 360
gtcagggagg ccgtgttgcc agacttgagg ccagagaagc gattagaaac ccctgagggc 420
cgatcagtga catcataaat catgagtttg ggggctttgc ctgggtgctg ttggtaccag 480
gagacatagt tataaaaacc aacgtcactg ctgggtccag tgcaggagat ggtgatcgac 540
tgt 543

<210> 971
<211> 416
<212> DNA
<213> Homo sapien

<400> 971
ccagactgac ttcaaaaaat taatgtgtat ccagggacat tttaaaaacc tgtacacagt 60
gtttatttgt gtttaggaagc aatttcccaa tgtacctata agaaatgtgc atcaagccag 120
cctgaccaac atgggtgaaac cccatctgta ctaaacataa aaaaatttagc ctggcatggt 180
ggtgtacgcc tgtaatccca gtgacttggg aggctgagggc aggagaatcg cttgaaccgc 240
ggaggcggag gttgcagtga gctaagatcg caccactgta ctccagcctg ggcaacagcg 300
agactccatc tcaaaaaaaaa aggaaatgtg tatcaagaac atgattatcc aggggtattt 360
totaattcag atcatcaaac tgattatata gaagagttag ctttaaaatg tttgca 416

<210> 972
<211> 242
<212> DNA
<213> Homo sapien

<400> 972
ccaaaaatcc caaaacatca ttttcaatca gtagagaagt gcttaggggtt gaaaattgat 60
ttcatttgct actgaatttg gtaaatcctg ggtaactttt atcaagatga agacatttta 120
ccctacctac tctagaaata tacaacaatg ttatatttta cactccttgg aaacatttga 180
ggaaaaaaat gcaatttgca cttcactttg ttggaatatc ccatagcact caataaactc 240
ag 242

<210> 973
<211> 347
<212> DNA
<213> Homo sapien

<400> 973
cctgcagggg atggaacctt ccagaagtgg ggcgctgtgg tggcgcttc tggagaggag 60
cagagatata cctgccatgt gcagcatgag ggtctgcca agccctcac cctgagatgg 120

```

gagctgtctt cccagccac catcccatc gtgggcatca ttgctggcct ggttctcctt 180
ggagctgtga tcaactggagc tgtggctcgt gccgtgatgt ggaggaggaa gagctcagga 240
cattttcttc ccacagatag aaaaggaggg agttacactc aggctgcaag cagtgcagct 300
gccagggcct ctgatgtgtc tctcacagct tgtaaagtgt gagacag 347

```

```

<210> 974
<211> 571
<212> DNA
<213> Homo sapien

```

```

<400> 974
gaaagagcga gatgcgagaa cacttttggc taaaaatctc ccttacaaag tcaactcagga 60
tgaattgaaa gaagtgtttg aagatgctgc ggagatcaga ttagtcagca aggatgggaa 120
aagtaaaggg attgcttata ttgaatttaa gacagaagct gatgcagaga aaacctttga 180
agaaaagcag ggaacagaga tcgatgggcg atctatttcc ctgtactata ctggagagaa 240
aggtcaaaat caagactata gaggtggaaa gaatagcact tggagtgggtg aatcaaaaac 300
tctggtttta agcaacctct cctacagtgc aacagaagaa actcttcagg aagtatttga 360
gaaagcaact tttatcaaaag taccacagaa ccaaaatggc aaatctaaag ggtatgcatt 420
tatagagttt gcttcattcg aagacgctaa agaagcttta aattcctgta ataaaaggga 480
aattgagggc agagcaatca ggctggagtt gcaaggaccc aggggatcac ctaatgccag 540
aagccagcca tccaaaactc tgtttgtcaa a 571

```

```

<210> 975
<211> 221
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(221)
<223> n = A,T,C or G

```

```

<400> 975
ctggaggtgc ctcanaaggt gcattctgct tcctgcaggg gcttgaaaca ccaaggcact 60
ccaggggatcc tggagtcaaa gcagcagccc cggttggtgc actccttggg ggtgacatgg 120
gggtagccgc agtccaccct gtccttggct ggcacggcac actggtttgc agacaggccc 180
acgtactcct cagcagagct ggaggacagc aaggccagga c 221

```

```

<210> 976
<211> 316
<212> DNA
<213> Homo sapien

```

```

<400> 976
ccatcagatt gtcacagact tttataaccc tttgatccct accaacgtta agtatgagtt 60
tggccctgcc atcttcattg gctgggcagg gtctgcccta gtcaccttg gaggtgcaact 120
gctctcctgt tcctgtcctg ggaatgagag caaggctggg taccgtgcac cccgctctta 180
ccctaagtc aactcttcca aggagtatgt gtgacctggg atctccttgc cccagcctga 240
caggctatgg gagtgtctag atgcctgaaa gggcctgggg ctgagctcag cctgtgggca 300
gggtgccgga caaagg 316

```

```

<210> 977
<211> 335
<212> DNA

```

<211> 550

<213> Homo sapien

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| ccatccccct | ttagaacgta | tcttaatgtg | aacataaatt | gttcttcatg | atgcttaaaa | 60 |
| gcttacatat | aattttcatt | cttagaaaaa | cgccacattt | tggatcctgg | atttttctga | 120 |
| atatcatgat | tgaaaaaaac | aaaacaaaaa | atgaacccaa | atcaaagtg | ggttaaactt | 180 |
| atatgagaaa | gatttttcaa | ccagatggtc | attcaaaaaa | gttgagctg | taagtgcg | 240 |
| cgactgagga | cacagggtta | attcctcgct | gctggtgaa | ggctagagaa | catcttcaa | 300 |
| agagggtagc | aagacgtgct | cctaggggag | gctcagtg | gtctcgtctg | cccaagcatt | 360 |
| ttcagtcctg | cttgggtcaat | gacatcgagt | aagtttttgg | catccacagc | cagggcgtga | 420 |
| gcagcagtc | gcatttgctt | tttgactctt | tgctggaggc | tggctatgac | atactgctgg | 480 |
| gccagtttca | tcttggtgat | gagctcacc | aggtcagagt | tcaatagctt | ctgtgccatc | 540 |
| tcaatctctc | | | | | | 550 |

<213> Homo sapien

| | | | | | | |
|-------------|-------------|------------|------------|-------------|-------------|-----|
| ccaaggtcag | aggctgatgc | aacaggccct | cttctcccca | gggccaggct | cctgtccagc | 60 |
| ctgggcactg | cccagagtga | tggcattggg | ccggatgctg | ttctgtctct | gcttggacac | 120 |
| cttcgcaaaag | atttctttca | ggacagtctc | aaaggctagc | tcaacattgg | tagagtccag | 180 |
| ggctgaggtc | tccaggaaga | gcagtcatt | gttttcagcg | aacattcggg | cctcctcagt | 240 |
| gggcacttcc | cgggcctggc | tgaggtcact | tttgttacc | acgagcatga | cgacgatcgt | 300 |
| ggcttcagca | tggtcataga | gctccttcag | ccatcgctcc | accacagcat | aggtctggtg | 360 |
| cttggttagg | tcaaacacca | ggagggcccc | cactgcacca | cgatagtacc | cttgaagaca | 420 |
| aagttataat | cttcctcagt | tccattcccc | atcttggtct | cgcattggagg | gtgcagggtgt | 480 |
| cttcggggac | agagggcgaca | aatctgtgtg | ttggctcaat | gccc | | 524 |

<213> Homo sapien

```

ccttcgtgccc ctaacagcca gtccctgttt aaagtggaag agacctgtgg ctgccgctgg      60
acctgccccct gtgtgtgcac aggcagctcc actcggcaca tcgtgacctt tgatgggcag      120
aatttcaagc  tgactggcag                                     140

```

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| tggagcggcc | gcccggcagg | tccaacgagt | cacaacagtg | caataggtag | aggattaaaa | 60 |
| actgcatcaa | acaggtgctg | aaaataaata | ctacctagga | gaaggagggtg | agagccctcg | 120 |
| tgtggggttt | gttttcgacc | ccttgagtg | gtgtgggggt | tgtcttcga | gccacgagcc | 180 |
| tggcctgtct | cgcggtgctg | ttcactctga | cagagtgcgc | ctgcagcacg | ttgcctccag | 240 |
| ggcccagcct | cccagaagcc | tcagagcatc | agagcatccg | tcccatcgga | tggaccagaa | 300 |
| acaagaaaat | gggggtgggt | gaatcacagc | tatcattcaa | aggaaaggaa | tttttttc | 358 |

<210> 985
 <211> 450
 <212> DNA
 <213> Homo sapien

<400> 985
 ctgaccccc tttgtccaca gctaagatgg cagcagaatg ctatgtcact atatacagaa 60
 acaagacaac ctgaagctaa atggatgccc cctgcagagt caacagggtcc agcctcacag 120
 tgcacgccct gagctacagc ctctcccaaa aggcattcttc cccacagcct caacgccgag 180
 caaggagcat caagggtttg tctcggttgt tttgttcttt ttacaaacta tagatatata 240
 cagttgaaaa ctcaggattt ctagccaata accatagtta ccaccacctt acaataaaaa 300
 agaaaatgcc agaaacatct ttaaattgcct tgtcacacca acagcaaagt gcacagagtg 360
 aggagaacac gagagtgcct tttcatttta aaaatgtttg gaaatatgta caactttgat 420
 acagtttcag ggtgctccag acacccatgg 450

<210> 986
 <211> 340
 <212> DNA
 <213> Homo sapien

<400> 986
 cctcctgcca gcagttcttg aagcttcttt ttcattcctg ctactctacc tgtattttctc 60
 agttgcagca ctgagtggtc aaaatacatt tctgggccac ctcagggaac ccatgcatct 120
 gcctggcatt taggcagcag agcccttgac cgtcccccac agggctctgc ctcacgtcct 180
 catctcattt ggctgtgtaa agaaatggga aaagggaaaa ggagagagca attgaggcag 240
 ttgaccatat ccagttttat ttattttatt ttaatttgtt tttttctcca agtccaccag 300
 tctctgaaat tagaacagta ggcggtatga gataatcagg 340

<210> 987
 <211> 227
 <212> DNA
 <213> Homo sapien

<400> 987
 ccaatgccc gagcaggccc tctttccatc ccgtgtcgga tgagctggtc aactatgtca 60
 acaaacggaa taccacgtgg caggccgggc acaacttcta caacgtggac atgagctact 120
 tgaagaggct atgtggtacc ttctgggtg ggcccaagcc accccagaga gttatgttta 180
 ccgaggacct gaagctgcct gcaagcttcg atgcacggga acaatgg 227

<210> 988
 <211> 241
 <212> DNA
 <213> Homo sapien

<400> 988
 cctcttttta ccagctccga ggtgattttc atattgaatt gcaaattcga agaagcagct 60
 tcaaacctgc cggggcttct cccgcctttt ttcccgcgcg cgggagaagt agattgaagc 120
 cagttgatta ggggtgcttag ctgttaacta agtgtttgtg ggtttaagtc ccattgggtct 180
 agtaagggtc tagcttaatt aaagtggctg atttgcgttc agttgatgca gagtggggtt 240
 t 241

<210> 989
 <211> 193
 <212> DNA

<213> Homo sapien

<400> 989

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccagccgtgt | cccagacttg | tagtttgatc | ttcttccct | ctatatccac | agtgcggatc | 60 |
| ttgaaatcaa | ttccgatgg | ggagatgtaa | gtgttggtga | agttgtctc | tgcaaagcga | 120 |
| atgatcagac | aagtcttgcc | cacccccgag | tccccgatca | gcagcaactt | gaagaggtgg | 180 |
| tcgtaggctt | tgg | | | | | 193 |

<210> 990

<211> 499

<212> DNA

<213> Homo sapien

<400> 990

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctcaaccaa | gagggttgat | ggcctccagt | caagaaactg | tggctcatgc | cagcagagct | 60 |
| ctctctcct | ccagcaggcg | ccatgcaagg | gcaggctaaa | agacctccag | tgcatcaaca | 120 |
| tccatctagc | agagagaaaa | ggggcactga | agcagctatg | tctgccaggg | gctaggggct | 180 |
| cccttgcala | cagcaatgct | acaataaagg | acacagaaat | gggggaggtg | ggggagccct | 240 |
| atthttataa | caaagtcaaa | cagatctgtg | cgttcattcc | cccagacaca | caagtagaaa | 300 |
| aaaaccaatg | ctgtggtttc | tgccaagatg | gaatattcct | cctcctagtt | ccacacatgg | 360 |
| cgtttgcaat | gctcgacagc | attgcactgg | gctgctgtct | ctgtgttctg | gcaccagtag | 420 |
| cttgggcccc | atatacactt | ctcagttccc | aacaaggggc | tatgggccga | ggggcaggct | 480 |
| ccaattttca | agcacacga | | | | | 499 |

<210> 991

<211> 262

<212> DNA

<213> Homo sapien

<400> 991

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgccagcca | ggctgtggtc | agtcctctgg | caggcaatct | tccggaccga | gagcctctgt | 60 |
| ccattagtgt | cagccccgag | ggggccacga | cggaggccgc | ccaatgtcca | ctgtgatatt | 120 |
| ggtgaagagt | ggttgccgag | acacctccaa | gacctggtac | cgcactgacc | caatgccgtc | 180 |
| ccgcttcatg | gtcagcttcg | tgttttgaat | cttggtaaac | ctctgagggg | taggttcgtt | 240 |
| atgcttctgc | cggctgtgct | tg | | | | 262 |

<210> 992

<211> 535

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(535)

<223> n = A,T,C or G

<400> 992

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ctgctgcttg | tgaaattcat | gtgtggtact | aagtaccta | catgaattat | ttcatttaac | 60 |
| cctcccaaca | gtctcctttg | tacgtgctgn | ncctctgcc | tggaaacact | gtttcccacc | 120 |
| cccaaccccc | aattcttctg | tttatttttc | ttgagacaga | gtctcactgt | gtagcccaga | 180 |
| ctggagtgca | gtggcgcgat | ctcggctcac | tccaatctcc | gcctcccggg | tccctgttca | 240 |
| agcagttctc | ctgcctcagc | ctcctgagta | gctgggatta | caggcacacg | ccaccatgtc | 300 |
| cagetaatth | ctgtattttt | agtagagatg | gggtttcacg | atgttggtta | ggatgggtctc | 360 |
| gatctctggt | cagagtcctt | tctgtaaata | tccttggtta | agaagcaatt | ttagactgta | 420 |

gctgttgcaa atgctttaag gaagaagcaa aacaactgtc agtcttnctg aaatgaagaa 480
actacaccag ggctgctata tcagagcaac cccaaccagc actncaatca tgatg 535

<210> 993
<211> 232
<212> DNA
<213> Homo sapien

<400> 993
ctgctgctct cccctcccag tctctactca ctgggatgag gttagggtcat gaggacacca 60
aaaacctaata aataaaca aaagccaaaca agccttagct tttcttaaag gctgaaatgc 120
ctggaagtgt ccttttattt ataaaataac ttttgtcata tttcttatac atgtttcttg 180
taagaaattc agaaactaca gacaaagaga gtggaaatta cccactgtca gg 232

<210> 994
<211> 203
<212> DNA
<213> Homo sapien

<400> 994
ccagcagatc atccacgacg accaccctct gtccctggctc cagggcgtct ttctgaatct 60
ccagctcagc cttcccgtac tccaggggaat aggaggccca cagagtgggg cctggcagct 120
tccccgcgtt tcggatgagc acgcagccca gtccaagctc ctggggccagg gaggggcca 180
agaggaagcc tcgggagtct agg 203

<210> 995
<211> 238
<212> DNA
<213> Homo sapien

<400> 995
ccatgcctgc cccgcccact ctgtatatat gtaagttaaa cccgggcagg ggctgtggcc 60
gtctttgtac tctgggtgatt tttaaaaatt gaatctttgt acttgcatg attgtataat 120
aattttgaga ccaggtctcg ctgtgttgct caggctggct ccaaactcct gagatcaagc 180
aatccgcccc cctcagcctc ccaaagtgtc gagatcacag gcgtgagcca ccaccagg 238

<210> 996
<211> 379
<212> DNA
<213> Homo sapien

<400> 996
ctgcagcctg ggactgaccg ggaggctctg accatttacc caccacaggt aggttgtgtt 60
ctgaacctca ggttcacagg tgaaggccac agcatccttg tcctccacgg ggttggaagt 120
gttgcctggag atggagggct tgggcagctc cgggtataca tggaactgtc cgttgcttc 180
ttcattcaca agatctgact ttatgacttg tagggatatag aatcctgtgt cattctgggt 240
gacgttctgg atcagcaggg atgcattggg gtatattgtc tctcgaccac tgtatgcggg 300
ccctggggta gcttgttgag ttcctattac atatcctaca attagactgt tgccatccac 360
tctttcgcct ttgtaccag 379

<210> 997
<211> 210
<212> DNA
<213> Homo sapien


```
<400> 997
ccatcccgaag caagattgca gatggcagt tgaagagaga agacatatc tacacttcaa      60
agcttttggtg caattccccat cgaccagagt tggtcgcacc agccttgaa aggtcactga    120
aaaatcttca attggattat gttgacctct accttattca tttccagtg tctgtaaagg     180
ccgtggagaa gtgtaaagat gcaggattgg                                     210

<210> 998
<211> 207
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)..(207)
<223> n = A,T,C or G

<400> 998
ggtggctgtg ctggggg cgc cccacaaccc tgctcccccg acgtccaccg tgatccacat      60
ncgcagcgcag acctccgtgc ccgaccatgt cgtctgggtcc ctgttcaaca ccctcttcat    120
gaacccctgc tgctgggct tcatagcatt cgctactcc gtgaagtcta gggacaggaa       180
gatggttggc gacgtgaccg gggccca                                         207

<210> 999
<211> 315
<212> DNA
<213> Homo sapien

<400> 999
ccaatgggct ttgctgtagc ttgctgaaat caccaagcag gagagattta accagaggcg      60
atgtgtccag tcaccagcat agagccatcc tctgtgtcac catccacacg cagggccttc     120
tggcagacct catgcaatgc cctccatggt aatattcatc agaaaatgga taattagggg     180
ggccagcaaa aatatcaagg gtcaaataat gcacatttct gtttaggccca tctatggcct     240
tcattctctc tgaagtcaac tggaattcaa acacctgcac gttctgtctg atgcgctgct     300
cattgtagct cttgg                                                         315

<210> 1000
<211> 186
<212> DNA
<213> Homo sapien

<400> 1000
ctgttactca agaagatgta tttaatgctt gacaataaga gaaaggaagt agttcacaaa      60
ataatagagt tgctgaatgt cactgaactt acccagaatg ccctgattaa tgatgaacta     120
gtggagtggg agcggagaca gcagagcgcc tgtattgggg ggccgcccaa tgcttgcttg     180
gatcag                                                                    186

<210> 1001
<211> 173
<212> DNA
<213> Homo sapien

<400> 1001
ccacaaaagcg gaaactcatt cacttttggc tttttccgcc ccagggtcaaa aatgcgaatc     60
```

```
<210> 1002
<211> 302
<212> DNA
<213> Homo sapien
```

```
<210> 1003
<211> 368
<212> DNA
<213> Homo sapien
```

```
<210> 1004
<211> 294
<212> DNA
<213> Homo sapien
```

```
<210> 1005
<211> 414
<212> DNA
<213> Homo sapien
```

| | | | | | | |
|-------------|-------------|------------|------------|-------------|------------|-----|
| <400> 1005 | | | | | | |
| ctgaagcact | cttcagagac | tacgtccaca | gacactgatg | ctgaggcctt | tcttgtaagt | 60 |
| gaagaaaaaag | gaatgcagca | aagaagagtt | cgacattgga | gtccttagtt | ccatcaggat | 120 |
| ccatttcgca | gccttttagca | tcatgtagaa | gcaaactgca | cctatggctg | agataggtgc | 180 |
| aatgacctac | aagattttgt | gttttctagc | tgtccaggaa | aagccatctt | cagtcttgct | 240 |
| gacagtcaaa | gagcaagtga | aaccatttcc | agcctaaact | acataaaaagc | agccgaacca | 300 |
| atgattaaaag | acctctaagg | ctccataatc | atcattaaat | atgccc aaac | tcattgtgac | 360 |

ttttttatattt atatacagga ttaaaatcaa cattaaatca tcttatttac atgg 414

<210> 1006

<211> 272

<212> DNA

<213> Homo sapien

<400> 1006

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ccggagccca | cggtgggcat | ggctgccaga | gcgctctgca | tgctggggct | ggtcctggcc | 60 |
| ttgtgtcct | ccagctctgc | tgaggagtag | gtgggcctgt | ctgcaaacca | gtgtgccgtg | 120 |
| ccagccaagg | acaggggtga | ctgcggctac | ccccatgtca | cccccaagga | gtgcaacaac | 180 |
| cggggctgct | gctttgactc | caggatccct | ggagtgcctt | gggtgtttcaa | gcccctgcag | 240 |
| gaagcagaat | gcaccttctg | aggcacctcc | ag | | | 272 |

<210> 1007

<211> 313

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (313)

<223> n = A,T,C or G

<400> 1007

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| cctgccttac | tctnttcctt | ttccccaggg | actcttggtt | ttcagaagcc | cctctggaat | 60 |
| gtectacctg | gcctaaccct | ataccagcag | tgcagacaag | gaggcactcc | tactatagtg | 120 |
| gggtccagccc | atggagagac | tcacttcctg | ccccaacacc | tcttccccta | gacctgagg | 180 |
| gccaggacaa | tgtcttagtg | ccttccaact | tggcagagtg | aggcccatg | agacagagag | 240 |
| aaagggggaa | gagggaata | cctttatcca | aataaatacc | catccaaaat | tatttgtgat | 300 |
| aggtgaaaaa | tgg | | | | | 313 |

<210> 1008

<211> 317

<212> DNA

<213> Homo sapien

<400> 1008

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctcaatgtc | gtgctagagg | ggccgaagaa | ggccgtgaac | gacgtgaatg | gcctgaagca | 60 |
| atgtttggca | gaattcaagc | gggatctgga | atgggttgaa | aggtctgatg | tgacactggg | 120 |
| tccggtaacc | gagatcggtg | gatctgaggc | gccagcacct | cagaacaagg | accagaaagc | 180 |
| tgttgatcca | gaagacgact | tccagcgaga | gatgagtttc | tatcgccaag | cccaggccgc | 240 |
| agtgtttgca | gtcttaccct | gcctccatca | gctcaaagtc | cctaccaagc | gacctactga | 300 |
| ttattttgcg | gaaatgg | | | | | 317 |

<210> 1009

<211> 456

<212> DNA

<213> Homo sapien

<400> 1009

| | | | | | | |
|------------|------------|-------------|-------------|------------|------------|-----|
| tttttttgta | gggtatagaa | aatacatttt | taattttgat | agagttcaca | aatgacagca | 60 |
| ttgacatttc | tttaaacaaa | tactttctgtc | aaggcacagc | attaccatgt | gtccccagat | 120 |
| gccaagagg | cagtgatctc | atgtccccct | gaggttttagc | agagccacca | atgtcaatag | 180 |

```

ggtggctgac ggggcctaga tttgctacca gataagccaa tgagacatgc tgtcagattt 240
atggttacat aatcaagtat ttaaaaagat gcacaatagg taactgcaat gagcttgctc 300
tgcatttagc gatagttcct ttcaaacaaa gaagatagtt ttcagtatca agaaggatgc 360
ctatatgtat gtcttccatg gagcctttcc tacaaattgc tttcattaca cattaagg 420
agttcagctt tattgtgacc ttcttgagtc attcag 456

```

<210> 1010

<211> 196

<212> DNA

<213> Homo sapien

<400> 1010

```

ctgggcatgg gctgaggaga ggtcttgctt gcccccttca actttccatc tcagaactat 60
aaactgctag gctgcaagga gagaagggct aagtgggggt cagacaggag agaagggcag 120
gaggcagtga gccccgatga cccaccaact ccaccaggcc ctgacaggga agcccccttg 180
gtagtatca ttttgg 196

```

<210> 1011

<211> 449

<212> DNA

<213> Homo sapien

<400> 1011

```

ccttgcggt gctgcgaaag gccacggcgc tgctgcccgc cggggccgag tactttgatg 60
gttcagagcc cgtgcagaac cgcgtgtaca agtcaactgaa ggtctgggtc atgctcgccg 120
acctgaagga gagcctcggc accttccagt ccaccaaggc cgtgtacgac cgcctcctgg 180
acctgcgtat cgcaacaccc cagatcgtca tcaactatgc catgttcttg gaggagcaca 240
agtacttcga ggagagcttc aaggcgtacg agcgcggcat ctgctgttc aagtggccca 300
acgtgtccga catctggagc acctacctga ccaaattcat tgcccgtat gggggccgca 360
agctggagcg ggcacgggac ctgtttgaac agyctctgga cggctgcccc ccaaaatatg 420
ccaagacctt gtacctgctg tatgcacag 449

```

<210> 1012

<211> 289

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (289)

<223> n = A,T,C or G

<400> 1012

```

ccaggaccac aacccccacgc ttagctggt agcgcagggc aatcagggct ggggttcgct 60
tgtgttttt tgccaaggca caaaggactg ggtcctccaa gagcaccggg gagttcgggt 120
ccacccatgg ttcttctcgg tgggatccca gagcactata ggcaaccaga acaatgtctt 180
ttgacttgca gaaatccagc agttttctct ggttgaagta aggatgacat tccacctggt 240
tgcagacagg ctgttacttg agccctggct tgnnaggat catctccag 289

```

<210> 1013

<211> 221

<212> DNA

<213> Homo sapien


```
<210> 1017
<211> 490
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|-------------|------------|------------|-------------|--|-----|
| <400> 1017 | | | | | | | |
| ctggaagaac | aaggcgaagt | tctgggtggct | gtctgcgatg | aatgtgccct | tggctttggc | | 60 |
| tgggtatgtc | acccggttag | ttttgggtgc | aatgctctga | tccttatcca | cgggtggaaag | | 120 |
| atcaacattt | gtgatgccaa | cttcagtggg | gatcttgact | ctgagctcta | cggatatttgc | | 180 |
| aatataccgg | ttgtcacctt | caacttcgac | aaggaagtca | taataaccac | tggaaaattt | | 240 |
| gacgttcatg | aaatttagtt | caaaaacatc | ccctacaggg | gtgaaggatg | tcttctggag | | 300 |
| gacagtggct | ctggaagcaa | cagatttagc | atgttctagt | ttaacagtgg | cctgagtcag | | 360 |
| aggctgagac | agaacattgg | tgacttgcaa | ccgcaagata | gcctgttcat | gagtgtcgga | | 420 |
| agcagancct | tcangcacia | ccacaactgg | cacgtggtag | cgattatgcg | agagcacagg | | 480 |
| cagacctcgg | | | | | | | 490 |

| | | | | | | | |
|-------------|-------------|-------------|------------|------------|------------|--|-----|
| <400> 1018 | | | | | | | |
| ggagtaagct | gagtacaagt | accatagcag | cagagctgca | aaaggtcttg | ggacctatag | | 60 |
| tcctaattgca | agataaggctc | atggggccta | aggccatggg | gcttgaggca | cccctagacc | | 120 |
| ctgagccttc | agcatttaag | ggagggtgtc | ccccattct | cgataggcca | tggtagacag | | 180 |
| atgggtctag | ccgaggtgct | ataactgctt | ggaccactgt | tgcagtccaa | cctagtactg | | 240 |
| acactatatg | gtttgaaacc | cgggtgtggac | aaagtagcca | atgggctgaa | cttagagcag | | 300 |
| tgtggatggg | gatcaccaag | gagggtgacac | tgatggtaat | ctgtatcaat | agctgggtgg | | 360 |
| tctaccaagg | cttaactttg | tggttaacta | cctggaaaat | acagaagttg | ctagtcggcc | | 420 |
| accaacccat | ttgggggtcaa | gccacgtggc | aagacctctg | ggaaatgggt | catcagaaac | | 480 |
| aggtaaccgt | ttatcatgtg | tca | | | | | 503 |

<400> 1019
cctgtgtatg gactagaggc ggggtgcacgg gtactgttcc tcacggcagt caagaggcccc 60

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aggetctgtg | ggetccagct | ctgcatttcc | cggttctggg | gttggggctg | ggatgacttc | 120 |
| ctgttggact | tgctgctggg | actggaactg | gaactgttcc | tcggagggcc | gaggagtcac | 180 |
| ctcttgataa | tcatagtagt | ctgggttgtc | gatctggtcg | ctatagtggg | tgtactggac | 240 |
| gtggtcaggg | aacggcggca | gcgggtccag | gtcatactgg | ccctgagcca | gcaagcctgc | 300 |
| aggcaggaat | agcaggaaga | ggtaggcagc | tctcatggca | acaaagag | | 348 |

<210> 1020

<211> 260

<212> DNA

<213> Homo sapien

<400> 1020

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccacacggcg | accgagggac | agatggggcc | ctgcgtccca | taggctgcct | gaaggtgggt | 60 |
| agggcggcct | gcggcatagt | ggggtggctg | tgggctccca | gcctggcccc | tgggaaccgt | 120 |
| gggagcacag | ggacaagcac | atggctatgg | aatgcagggg | gacccaagga | caagcgagtt | 180 |
| gcggggatct | ctactgtgac | catgcagaat | tgatcgcagt | ctgctgcgcc | accaccacct | 240 |
| catgttcccg | aggggaacag | | | | | 260 |

<210> 1021

<211> 407

<212> DNA

<213> Homo sapien

<400> 1021

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ccttatgact | ataacggccc | acgagaaaaa | tatggaatcg | ttgattacat | gatcgagcag | 60 |
| tccgggcctc | cctccaagga | gattctgacc | ctgaagcagg | tccaggagtt | cctgaaggat | 120 |
| ggagacgatg | tcatcatcat | cggggtcttt | aagggggaga | gtgaccacagc | ctaccagcaa | 180 |
| taccaggatg | ccgctaacaa | cctgagagaa | gattacaaat | ttcaccacac | tttcagcaca | 240 |
| gaaatagcaa | agttcttgaa | agtctcccag | gggcagttgg | ttgtaatgca | gcctgagaaa | 300 |
| ttccagtcca | agtatgagcc | ccggagccac | atgatggacg | tccagggctc | cacccaggac | 360 |
| tcggccatca | aggacttcgt | gctgaagtac | gccctgcccc | tggttgg | | 407 |

<210> 1022

<211> 140

<212> DNA

<213> Homo sapien

<400> 1022

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccaccccaga | gtgggagagg | ctgggagggt | gggaggctgt | ggagagaagt | gagcaagggtg | 60 |
| ctcttgaacc | tgtgtcatt | ttgcaat | atcagtaatt | tgacttagag | tttttacgaa | 120 |
| acctcttttg | ttgtccttgc | | | | | 140 |

<210> 1023

<211> 280

<212> DNA

<213> Homo sapien

<400> 1023

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctggaggtgc | ctcagaagg | gcattctgct | tcctgcaggg | gcttgaaaca | ccaaggcact | 60 |
| ccagggatcc | tggagtcaaa | gcagcagccc | cggttgttgc | actccttggg | ggtgacatgg | 120 |
| gggtagccgc | agtccaccct | gtccttggct | ggcacggcac | actggtttgc | agacaggccc | 180 |
| gcgtactcct | cagcagagct | ggaggacagc | aaggccagga | ccagccccag | catgcagagc | 240 |
| gctctggcag | ccatgaccac | cgtgggctcc | gggacgcagc | | | 280 |

<210> 1024
 <211> 274
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(274)
 <223> n = A,T,C or G

<400> 1024
 cctggctgag caggcagagc accctgggac cccagggcag aaggaccct gccctccagt 60
 cccaagacc caggcccgtc tccactcata cagccacct acatgtgacg tcagccctga 120
 aaaggtaaca ggaaagtcca gaacaaaaaac aaaaccccaa aagtaaaaag gctacgtgta 180
 gcagagtaat accggaaacg ttatatacac aggcgggtgat ggccccctcg gaagtgtccg 240
 ggtcacttag ggggcactgc anaggtccct gtgg 274

<210> 1025
 <211> 446
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(446)
 <223> n = A,T,C or G

<400> 1025
 gcaaagagtg tactgtgctt gaggcagagc actcacacat aaatggctgt gtgtggaatt 60
 gcttgccaaa gaagtttcta gcctttccct tccccctaac tgcacaggg aagaattott 120
 atctctagct tggtttccac atgaggtttt tctgagaagg gcttgggaca agaagtctgt 180
 catgttagtt aagcaggcaa gaaatcctac taatccagtt ttgtttgaaa gttgtttgtc 240
 cgtatgattt tttaaaagtc aagtttaatt tcaaaaaacc ttttttttct gagattactt 300
 ttggggtaat atttaaaatg agagacattt tgtaaccctg taaaatacat aggggaatata 360
 acattccagt gtatacaaag aaggcaaatt cttaaatcaa ataaagcgca ttataaaatc 420
 aaaaaanaaa naaaaaaaaaa aaaaaa 446

<210> 1026
 <211> 189
 <212> DNA
 <213> Homo sapien

<400> 1026
 ctgtgagaga gatgctcaat atgccccagg ctatgacaaa gtcaaggaca tctcagaggt 60
 ggtcacccct cgggttccttt gtactggagg agtgagtccc tatgctgacc ccaatacttg 120
 cagaggtgat tctggcggcc ccttgatagt tcacaagaga agtcgtttca ttcaagttgg 180
 tgtaatcag 189

<210> 1027
 <211> 92
 <212> DNA
 <213> Homo sapien

<400> 1027

ccagaccctc cttagtagacag gatctcggac cacaaaccaa ggagtctcgt ggccttggat 60
tcccagaccc taggatggta tccctctgac ag 92

<210> 1028
<211> 438
<212> DNA
<213> Homo sapien

<400> 1028
ctgaaaagcc atcttttgcac tgttccctcat ccgcctcctt gctcgccgca gccgcctccg 60
ccgcgcgcct cctccgccgc cgcggactcc ggcagcttta tcgccagagt ccctgaactc 120
tcgcttttctt tttaatcccc tgcatcggat caccggcggtg ccccaccatg tcagacgcag 180
ccgtagacac cagctccgaa atcaccacca aggacttaaa ggagaagaag gaagttgttg 240
aagaggcaga aaatggaaga gacgcccctg ctaacgggaa tgctaataag gaaaatgggg 300
agcaggaggc tgacaatgag gtagacgaag aagaggaaga aggtggggag gaagaggagg 360
aggaagaaga aggtgatggg gaggaagagg atggagatga agatgaggaa gctgagtcag 420
ctacggggcaa gcgggcag 438

<210> 1029
<211> 330
<212> DNA
<213> Homo sapien

<400> 1029
ccagccgcat gggagtggag gcagtcacgc ccttgctaga ggccaccccg gacacccag 60
cttgcgctcgt gtcactgaac gggaaccacg ccgtgcgcct gccgctgatg gagtgcgtgc 120
agatgactca ggatgtgcag aaggcgatgg acgagaggag atttcaagat gcggttcgac 180
tccgagggag gagctttgcg ggcaacctga acacctacaa gcgacttgcc atcaagctgc 240
cggatgatca gatcccaaag accaatcgca acgtagctgt catcaacgtg ggggcacccg 300
cggctgggat gaacgcggcc gtacgctcag 330

<210> 1030
<211> 228
<212> DNA
<213> Homo sapien

<400> 1030
ctggagactc tgggccagga gaagctgaag ctggaggcgg agcttgccaa catgcagggg 60
ctgggtggagg acttcaagaa caagtatgag gatgagatca ataagcgtac agagatggag 120
aacgaatttg tctcatcaa gaaggatgtg gatgaagctt acatgaacaa ggtagagctg 180
gagtctcgcc tggaagggct gaccgacgag atcaacttcc tcaggcag 228

<210> 1031
<211> 294
<212> DNA
<213> Homo sapien

<400> 1031
ccacaaagcc attgtatgta gcttttagctc agcgcaaaga agagcgccag gctcacctca 60
ctaaccagta tatgcagaga atggcaagtg tacgagctgt gcccaccct gtaatcaacc 120
cctaccagcc agcacctcct tcaggttact tcatggcagc tatccacag actcagaacc 180
gtgctgcata ctatcctcct agccaaattg ctcaactaag accaagtccc cgctggactg 240
ctcaggggtgc cagacctcat ccattccaaa atatgcccgg tgctatccgc ccag 294

<210> 1032
 <211> 278
 <212> DNA
 <213> Homo sapien

<400> 1032
 ggaggtatta cagacagcac tgcactttgg agttgggcag ctacatcgag gacctctttg 60
 tgggtccacag tgacctctcc agcattgtga tcttgataa ctccccaggg gcttacagga 120
 gccatccaga caatgccatc cccatcaaat cctgggttcag tgacccagc gacacagccc 180
 ttctcaacct gctcccaatg ctgggtgccc tcagggttcac cgctgatgtt cgttcogtgc 240
 tgagccgaaa ccttcaccaa catcggtctt ggtgacgg 278

<210> 1033
 <211> 155
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(155)
 <223> n = A,T,C or G

<400> 1033
 cgcgttcanc catgttnaaa ccgattgcat naacttcgaa accggcccgc ccgcccggcg 60
 ctggagaggg gcanngggag aagcagagag tttatcattc atctgtacac atagacgttt 120
 cttctttaaa taacaccacg ggcgggagcc ccac 155

<210> 1034
 <211> 401
 <212> DNA
 <213> Homo sapien

<400> 1034
 ctggaccagc acccattga cgggtacctc tcccacaccg agctggctcc actgcgtgct 60
 cccctcatcc ccatggagca ttgcaccacc cgctttttcg agacctgtga cctggacaat 120
 gacaagtaca tcgccctgga tgagtgggcc ggctgcttcg gcatcaagca gaaggatctc 180
 gacaaggatc ttgtgatcta aatccactcc tcccacagta ccgattctc tctttaaccc 240
 tccccttcgt gtttccccca atgtttaaaa tgtttgatg gtttggtgtt ctgcctggag 300
 acaagggtgct aacatagatt taagtgaata cattaacggt gctaaaaatg aaaattctaa 360
 cccaagacat gacattctta gctgtaactt aactattaag g 401

<210> 1035
 <211> 333
 <212> DNA
 <213> Homo sapien

<400> 1035
 ctgagctggg ggttgaatth ctccaggcac tccctggaga gaggaccag tgacttgtcc 60
 aagtttacac acgacactaa tctcccctgg ggaggaagcg ggaagccagc caggttgaac 120
 tgtagcgagg cccccaggcc gccaggaatg gaccatgcag atcactgtca gtggagggaa 180
 gctgctgact gtgattaggt gctggggtct tagcgtccag cgcagcccgg gggcatcctg 240
 gaggtctgct tccttagggc atggtagtca ccgcgaagcc gggcaccgtc ccacagcatc 300
 tcctagaagc agccggcaca ggaggggaagg tgg 333

<210> 1036
 <211> 198
 <212> DNA
 <213> Homo sapien

| | |
|--|-----|
| <400> 1036 | |
| ccaatgtaca tgggtggacta tgccgggctg aacgtgcagc tcccgggacc tcttaattac | 60 |
| tagacctcag tactgaatca ggacctcact cagaaagact aaaggaaatg taatttatgt | 120 |
| acaaaatgta tattcggata tgtatcgatg ccttttagtt tttccaatga tttttacact | 180 |
| atattcctgc caccaagg | 198 |

<210> 1037
 <211> 289
 <212> DNA
 <213> Homo sapien

| | |
|---|-----|
| <400> 1037 | |
| ctggagatga tcctcaacaa gccagggtc aagtacaagc ctgtctgcaa ccagggtggaa | 60 |
| tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattggt | 120 |
| ctgggtgcct atagtgtctt gggatccac cgagaagaac catgggtgga cccgaactcc | 180 |
| ccggtgctct tggaggacc agtcctttgt gccttggcaa aaaagcacia gcgaaccca | 240 |
| gccctgattg ccctgcgcta ccagctacag cgtgggggtg tggctcctgg | 289 |

<210> 1038
 <211> 368
 <212> DNA
 <213> Homo sapien

| | |
|--|-----|
| <400> 1038 | |
| ccagacgtgg tggctcacac ctgcaatccc agcaccttag gaggcagagg caggaggatc | 60 |
| cttgagggtca ggagttcgag accagcctcg ccaacatggt gaaaccccat ttctactaaa | 120 |
| aatacaaaaa attagccaag tgtggtggca tatgcctgta atcccaacta ctcagaaggc | 180 |
| cgaggcagga gaattacttg aacgcaggag aatcactgca gcccaggagg cagaggttgc | 240 |
| agtgagccga gattgcacca ctgcactcca gcctgggtga cagagcaaga ctccatctca | 300 |
| gtaaataaat aaataaataa aaagcgctgc agtagctgtg gcctcaccct gaagtcagcg | 360 |
| ggccccagg | 368 |

<210> 1039
 <211> 417
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(417)
 <223> n = A,T,C or G

| | |
|---|-----|
| <400> 1039 | |
| ctgggcctat gctgggtcatg aacgggtcctg gaaaatgact cccttccttc agtatctgca | 60 |
| tcctcatgaa gtcattcatt ttggagatcg tgtcttcact tttcttggtg aagaaactgc | 120 |
| tggatggagt tgttggtggc atctgaggag tccgaagatg gctctcaggg aaggttgtgc | 180 |
| tggcctctga aggatttgga agctgactct gttcctgggg tagctnnatg ctcttgggg | 240 |
| cattgnttct cgggtttgnt tttttcttta tctggataaa actatgcatt tctgaaatca | 300 |
| gttttgacat ctggttcttt tttcctaagt cgaaagcaga aaagttggaa gcttatctcc | 360 |

ttcttcacag ggggatattg tggacattgn nctgtcccca ctacatccat ttttcct 417

<210> 1040

<211> 409

<212> DNA

<213> Homo sapien

<400> 1040

| | | | | | | |
|------------|------------|-------------|-------------|------------|------------|-----|
| ctgtccaatg | gcaacaggac | cctcactcca | ttcaatgtca | caagaaatga | cgcaagagcc | 60 |
| tatgtatgtg | gaatccagaa | ctcagtgagt | gcaaaccgca | gtgaccaggt | caccctggat | 120 |
| gtcctctatg | ggccggacac | ccccatcatt | tccccccag | actcgtctta | cctttcggga | 180 |
| gcgaacctca | acctctcctg | ccactcggcc | tctaaccocat | ccccgcagta | ttcttggcgt | 240 |
| atcaatggga | taccgcagca | acacacacaa | gttctcttta | tcgccaaaat | cacgccaaat | 300 |
| aataacggga | cctatgcctg | ttttgtctct | aacttggcta | ctggccgcaa | taattccata | 360 |
| gtcaagagca | tcacagtctc | tgcattctgga | acttctcctg | gtctctcag | | 409 |

<210> 1041

<211> 492

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(492)

<223> n = A,T,C or G

<400> 1041

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| cctcggtccc | acacctccgc | tgtgaccaca | gcctcaggtc | aagctgtgct | ggggccatcc | 60 |
| accttccttt | gccatttaga | agatggggct | tggagcttgg | caacacagaa | attgacatca | 120 |
| gccttataaa | accttggctg | aacctaccga | cctccaggag | aatttcagcc | aaaacaaaaa | 180 |
| agcaaataca | cagagggacc | ctggaaccag | aatccctccc | catgggaaaag | acgaaggcac | 240 |
| agagattcga | gccaagtttc | ccaacatggt | ggtgtttgca | gaaaagtccg | gtcacgtcac | 300 |
| acacagcaca | gaggcaagaa | gcgaaggcag | tggcattcac | aggactactt | tatattaaag | 360 |
| tttattacat | ttggaaaatc | tactgtacag | ggaaaaaccc | attggattaa | gtagagtttt | 420 |
| gccaaaagca | aaagactatc | actctttgga | aaatattcct | gattccagcc | cangggcccag | 480 |
| ggtggggcca | ca | | | | | 492 |

<210> 1042

<211> 125

<212> DNA

<213> Homo sapien

<400> 1042

| | | | | | | |
|------------|-----------|------------|-------------|------------|------------|-----|
| cctggctctg | atccagtgc | ccctctcacc | aaagaactcg | gtttaaccag | ggctctgtaa | 60 |
| gacctctccc | accagagac | ttgtgtggcc | tgggtgtggcc | tgtgtgtcgg | attccttcct | 120 |
| gtcag | | | | | | 125 |

<210> 1043

<211> 459

<212> DNA

<213> Homo sapien

<400> 1043

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|----|
| ccagcctgga | gataaggggtg | aaggtggtgc | ccccggactt | ccaggtatag | ctggacctcg | 60 |
|------------|-------------|------------|------------|------------|------------|----|

```
<210> 1044
<211> 368
<212> DNA
<213> Homo sapien
```

```
<210> 1045
<211> 315
<212> DNA
<213> Homo sapien
```

```
<210> 1046
<211> 317
<212> DNA
<213> Homo sapien
```

```
<210> 1047
<211> 412
<212> DNA
<213> Homo sapien
```

$\langle 220 \rangle$

```
<400> 1050
ctgcagcctg ggactgaccg ggaggctctg attattttacc caccacaggt aggttgtggt      60
ctgaatctca ggttcacagg ttaaggctac agcatcctca tctccacgg ggttggagtt    120
gttgctggtg atgaagggtt tgggtggctc tgcatagact gtgatcgtcg tgactgtggt    180
```

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| cctattgagg | ccagtgtctg | agttatgggc | ttggcacgta | taggatccac | tattattcac | 240 |
| agtgatgttg | gggataaaga | gctcttgggt | ggattgctgg | aaagtcccat | tgacaaacca | 300 |
| agagtactgt | gcaggtgggt | tagaggctgc | gtggcaggag | aggttcagat | tttccctga | 360 |
| tctgtaagat | gtgttttagag | gggaaatggg | gggggcatcc | gggccataga | ggacattcag | 420 |
| gatgactgaa | tcactgcgcc | tggcactcac | tgggttctgg | gtttcacatt | tg | 472 |

<210> 1051

<211> 249

<212> DNA

<213> Homo sapien

<400> 1051

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaccaaccg | tggcatcacg | cgaatccggg | gcaccagcta | ccagagccct | cacggcatcc | 60 |
| ccatagacct | gctggaccgg | ctgcttatcg | tctccaccac | cccctacagc | gagaaagaca | 120 |
| cgaagcagat | cctccgcata | cgggtgcgag | aagaagatgt | ggagatgagt | gaggacgcct | 180 |
| acacggtgct | gacccgcata | gggctggaga | cgctactgcg | ctacgccata | cagctcatca | 240 |
| cagacctgc | | | | | | 249 |

<210> 1052

<211> 289

<212> DNA

<213> Homo sapien

<400> 1052

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaggaccac | aacccccacg | tgtagctggt | agcgcagggc | aatcagggct | ggggttcgct | 60 |
| tgtgcttttt | tgccaaggca | caaaggactg | ggctctccaa | gagcaccggg | gagttcgggt | 120 |
| ccacccatcg | tttgtctcgt | tgagatccca | gagcactata | ggcaaccaga | acaatatctt | 180 |
| tgcacttgca | gaaatctagc | aatttactcc | ggttgaaata | cggatgacat | tctacctggt | 240 |
| tgcagacagg | cttgtaactg | agtcttggtc | tggtgaggat | catctccag | | 289 |

<210> 1053

<211> 199

<212> DNA

<213> Homo sapien

<400> 1053

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccacgactgc | atgcccgcgc | ccgccagggt | atacctccgc | cggtgaccca | ggggctctgc | 60 |
| gacacaagga | gtctgcatgt | ctaagtgcta | gacatgctca | gctttgtgga | tacgcggact | 120 |
| ttgttgctgc | ttgcagtaac | cttatgccta | gcaacatgcc | aatctttaca | agaggaaacc | 180 |
| gtaagaaagg | gcccagccg | | | | | 199 |

<210> 1054

<211> 224

<212> DNA

<213> Homo sapien

<400> 1054

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tgcacctgt | gaagcaggag | acagatgctg | cattttcact | gttgtttgtc | ctctgttttt | 60 |
| gtagcatccc | cgggaacttc | cccatcagcc | aggggcttgt | ccccaccacc | cttcacctgg | 120 |
| ctttccagtt | ggctgagacg | ctgcttcata | ttcatctggg | tggcgttgta | ctcagccagg | 180 |
| aggcgtgcaa | acctggtctg | cagggcgctc | agggaggacc | ccag | | 224 |

<210> 1055

<211> 390

<212> DNA
<213> Homo sapien

<400> 1055
cctcttatta gggctctggt agcggcgggcg gcggaccctt ggggtctgga cgcaacggcg 60
gcgggagcat gaacgcccct ccagccttcg agtcgttctt gctcttcgag ggcgagaaga 120
agatcaccat taacaaggac accaagggtac ccaatgcctg tttattcacc atcaacaaag 180
aagaccacac actgggaaac atcattaaat cacaactcct aaaagaccg caagtgtat 240
ttgtctggcta caaagtcccc cacccttgg agcacaagat catcatccga gtgcagacca 300
cgccggacta cagccccag gaagcctttg ccaacgccat caccgacctc atcagtgagc 360
tgtccctgct ggaggagcgc tttcgggtgg 390

<210> 1056
<211> 450
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G

<400> 1056
ccagcatcac cttttggtcc nnacactcca gggctgccag gagcaccagt gttacccgca 60
ggacctgggg gccatcctt gcctggagaa ccgctgggac ctgggggtcc tgggttacca 120
ttactaccag gaggaccagg aagaccacga gcaccaggga agccagcagc accagggtcca 180
ccaggactgc caggttcacc tttgacacct tggggaccag gaggaccagn angtccagaa 240
cctccagggg gtcttgcaac tccaggaggg cctccttcac ctttctcacc cggagcccct 300
ctttctcctt taccaccagg ttcaccattc tgtccaggag caccagggaa accagcaggt 360
cctggagggc cagtttnacc tctctcacca nggctaccac gaggtccagc tatacctgga 420
agtccggggg caccaccttc acccttacct 450

<210> 1057
<211> 337
<212> DNA
<213> Homo sapien

<400> 1057
tgagcgggcg cccggcaggt cctcgcctgg agggccccgg gcagcacagg gaggacgagc 60
ttgtccagca gagggtctgg cagaggttcc cgcagaggtt tgggcagggg gtctgacatc 120
cctggctcct gctctggctc tggctgccgg gatttgcaca ggcccaggtg catacagatg 180
ccgtttgagt caatctggtt ctggaagtag tcgatgacca gggggaagta gtcgtcaagc 240
acttggttgc actggggcat gagcagcttc aaggggagga cgttgactc ctgctccagg 300
aacttctca tcgtgtcctg gaaaatggcc tccttgg 337

<210> 1058
<211> 237
<212> DNA
<213> Homo sapien

<400> 1058
ctgggggactg ggaatgctag catatggtat ctcaagttgg ctctcagaac taaacgggga 60
taagggccta gaatggaaga gggaaccagc cagacctca gtccttcctg tcctggactg 120
ggagccacag atgtccctgt gatctgtcac tgccctgatc tgggtcttca gccattaaa 180

ctcagtgtca tcttcagtca ccaacggggg tcttggtgtc cttccaaacc cctttgg

237

<210> 1059

<211> 210

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(210)

<223> n = A,T,C or G

<400> 1059

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| agcccatccc | cccggtccc | tcctagtctg | ccctgcgccc | tctgtccccg | ggtttcagag | 60 |
| acaacttccc | aaagcacaaa | gcagtttttc | cccctagggg | tgggaggaag | caaaagactc | 120 |
| tgtacctact | ttgtatgtgt | ataataattt | gagatgtttt | taattattnn | gattgctgga | 180 |
| ataaagcatg | tggaatgac | caaaaaaaaa | | | | 210 |

<210> 1060

<211> 564

<212> DNA

<213> Homo sapien

<400> 1060

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctggccacag | agcccagcaa | gtccttcctg | ggagagaaga | gttagggctg | atactgaagg | 60 |
| tctctttcac | atctgggcac | acgtctgcct | tcaggctgta | agaatttcat | ttgtcgattg | 120 |
| ttaaataaaa | ccaggagaaa | gcaatgcagg | tctctgggaa | tctcatccct | tccataagga | 180 |
| aaatgctctg | ccaattcaag | tttcattcag | tcaggaagac | agaaggattt | aaggcttcgg | 240 |
| tgacaattat | aatcctctga | gaaattattt | ccccttaaag | tcaagataag | ataatagtgt | 300 |
| ttactgtact | ttctcttgac | tcttgaaatc | cctggtattg | ggtgtaggca | acttgcacct | 360 |
| gcaatgaagt | cgcaggaga | ggaaggtctc | tcctcccccg | aaagctatcc | caggtcacat | 420 |
| gcgtggcgaa | tgccactga | acctcggctc | tcatggaagc | aggaaagaca | ccgagattca | 480 |
| agccttctag | taggttgagg | acgctgtgct | catggcatct | tcggagattt | tggtactggc | 540 |
| aggggtggat | gcttgcaaaa | tact | | | | 564 |

<210> 1061

<211> 267

<212> DNA

<213> Homo sapien

<400> 1061

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| cctatggagg | tgccatgat | gtcatgagct | ctaagcacct | tttggtgat | accaactatg | 60 |
| cctggccac | cgagagatt | gcggtcatgg | gagcaaaggg | cgctgtggag | atcatcttca | 120 |
| aaggggcatga | gaatgtggaa | gctgctcagg | cagagtacat | cgagaagttt | gccaaacctt | 180 |
| tcctgcagc | agtgcgagg | tttgtggatg | acatcatcca | accttcttcc | acacgtgccc | 240 |
| gaatctgctg | tgacctggat | gtcttgg | | | | 267 |

<210> 1062

<211> 603

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<223> n = A, T, C or G

| | | | | | | |
|------------|-------------|------------|------------|-------------|-------------|-----|
| ctggtcatct | tgctcatgtga | agaccatctt | cctacagagt | ctaggtctggc | cgtcgttgaa | 60 |
| gtcctcacca | gtactacacc | acttttcttc | accaaccccc | atcctattct | tgagttgcag | 120 |
| gatacacttg | ctctctggaa | gtgtgtcctt | acccttctgc | agagtgagga | gcaagctggt | 180 |
| agagatgcag | ccacggaaac | cgtgacaact | gccatgtcac | aagaaaatac | ctgccagtca | 240 |
| acagagtttg | ccttctgcca | ggtggatgcc | tccatcgctc | tggccctggc | cctggccgtc | 300 |
| ctgtgtgatc | tgctccagca | gtgggaccag | ttggcccttg | gactgcccat | cctgctggga | 360 |
| tggctgtttg | gagagagtga | tgacctcgtg | gcctgtgttg | agagcatgca | tcaggttggaa | 420 |
| gaagactacc | tgtttgaaaa | agcagaagtc | aacttttggg | ccgagacctt | gatctttgtg | 480 |
| aaatacctct | gcaagcacct | cttctgtctc | ctctcaaaag | tccggtctggc | gtncoccaaag | 540 |
| ccctgagatg | ctctgtcacc | ttcaaaggat | ggtgtcagag | cagtgccacc | tnctgtctca | 600 |
| qtt | | | | | | 603 |

<211> 222

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccatcgtgga | tcactgagat | gcagtggcgg | tcccgtagc | tggcccgtgg | catgccccc | 60 |
| tggaagatgg | tgaagggcaa | cccctgccta | gtggtcagcc | ggaggattct | ggtaatcgt | 120 |
| ctgcaaggaa | agggaccgta | aggcacgagg | ctgcggaggg | gctctggttg | ctgggcttcg | 180 |
| ctggacacgg | gccactggca | gtagctgccg | tcagagtgc | ag | | 222 |

<211> 72

<213> Homo sapien

<221> misc feature

<223> n = A, T, C or G

gatgatcaat atnnaactgga acacatgcat gcttttggaa tgtataatta cctgcactgt 60
gattcatggt at 72

<211> 251

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| gtggccgtga | tggatagcga | caccacaggc | aagctgggct | ttgaggaatt | caagtacttg | 60 |
| tgaacaaca | tcaaaaggtg | gcaggccata | tacaaacagt | tcgacactga | ccgatcaggg | 120 |
| accatttgc | gtagtgaact | cccaggtgcc | tttgaggcag | cagggttcca | cctgaatgag | 180 |
| catctctata | acatgatcat | ccgacgctac | tcagatgaaa | gtgggaacat | ggattttgac | 240 |
| aacttcatca | g | | | | | 251 |

<210> 1066

<211> 289
 <212> DNA
 <213> Homo sapien

<400> 1066
 ctggagatga tcctcaacaa gccagggctc aagtacaagc ctgtctgcaa ccaggtggaa 60
 tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattggt 120
 ctggttgctt atagtgtctt gggatccac cgagaagaac catgggtgga cccgaactcc 180
 ccagtgtctt tggaggaccc agtcctttgt gccttggcaa aaaagcacia gcgaacccca 240
 gccctgattg cctgcgcta ccagctacag cgtgggggtg tggctctgg 289

<210> 1067
 <211> 301
 <212> DNA
 <213> Homo sapien

<400> 1067
 ctgtagttag ctgaagtcgc taaacaggac ggatttaagt agaggtgata tgtccagtca 60
 ccggcataga gacgtcctt gcgtcaccat ccacacacag ggcttctggt agacatcagg 120
 caaagctctc catgttaata ttcattctgaa tatggataat taggggtggct agcaaaacta 180
 tcaactgtta aatagtggag atttctgtct aggccatcta tggctttcat gtctccgca 240
 gtcaactgga actcaaaaac ctgcacgttc tgtctgatgc gctgctcatt gtagctcttg 300
 g 301

<210> 1068
 <211> 255
 <212> DNA
 <213> Homo sapien

<400> 1068
 ccagcagttc ctctttgcct tatatttggt gtacgcccgg ccagccttca agatggggtt 60
 gtcaattcgg ccacctccag ccaccacacc aaccacagct ctggttgctg aggagataac 120
 cttcttgagg ccggagggca gttcacacg ggtcttcttg gtctcagggt tgtgggagat 180
 aacggtggca tagttccctg atgcccgggc cagcttgcca cggctctccag gttctctctc 240
 caggcagcac acgat 255

<210> 1069
 <211> 77
 <212> DNA
 <213> Homo sapien

<400> 1069
 ctggacaggc tccagcaccg gcccaaacac gcccagacct cggcaggcac cacctgggtc 60
 tcccaccag aaagtcc 77

<210> 1070
 <211> 163
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(163)
 <223> n = A,T,C or G

```
<210> 1071
<211> 246
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|------------|------------|-------------|-------------|--|-----|
| <400> | 1071 | | | | | | |
| ctgaccggac | cggncatgcc | cgtccggaac | gtctataaga | aggagaaagc | tcgagtcatc | | 60 |
| actgaggaag | agaagaattt | caaagccttc | gctagtctcc | gtatggcccg | tgccaacgcc | | 120 |
| cggctcttcg | gcatacgggc | aaaaagagcc | aaggaagccg | cagaacagga | tgttgaaaaa | | 180 |
| aaaaaaaaaa | gccctcctgg | ggacttgga | tcagtcggca | gacaaaaaaaa | aaaaaaaaaaa | | 240 |
| aacaaa | | | | | | | 246 |

```
<220>
<221> misc_feature
<222> (1) ... (224)
<223> n = A,T,C or G
```

```
<210> 1073
<211> 301
<212> DNA
<213> Homo sapien
```

<210> 1074
<211> 132

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(132)
<223> n = A,T,C or G

| | |
|--|-----|
| <400> 1074 | |
| caagcttttt tttttttttt tttttttttt ttcgctcaaa nactttnttt tattantaca | 60 |
| tgggctggna ttgatggnaa gggacaaatg tanttggcaa ccatgggttag catcggatgc | 120 |
| ccatcccaat gg | 132 |

<210> 1075
<211> 301
<212> DNA
<213> Homo sapien

| | |
|---|-----|
| <400> 1075 | |
| ctgtagttga ctgaagtcgc taaacaggac ggattttaagt agaggtgata tgtccagtca | 60 |
| ccggcataga gacgtcctct gcgtcaccat ccacacacag ggcttctggg agacatcagg | 120 |
| caaagctctc catgttaata ttcattctgaa tatggataat taggggtggc agcaaaaacta | 180 |
| tcactgttaa aatagtgagg atttctgtct aggccatcta tggctttcat gtccctctgca | 240 |
| gtcaactgga actcaaaaac ctgcacgttc tgtctgatgc gctgctcatt gtagctcttg | 300 |
| g | 301 |

<210> 1076
<211> 436
<212> DNA
<213> Homo sapien

| | |
|--|-----|
| <400> 1076 | |
| ctgctgggat gaatgccaa tttttcagcc ataaggtagc gaaatctagc agaatccaga | 60 |
| ttacatccac ttccaatcac gcgggtgtttg ggtaatccac ctagtctcca ggtaacatac | 120 |
| gtaagaatgt ccactgggtt ggaaaccaca attatgatgc aatcaggact gtacttgacg | 180 |
| atctgaggaa taatgaattt gaagacatta acatttctct gcaccagatt gagccgactc | 240 |
| tcccttctct gctgacggac tcctgcagtt actactacaa tcttagaatt ggcggtcaca | 300 |
| gaataatctt tatctgccac aatttttaggt gtctgaagaa ataagctccc atgctgcaga | 360 |
| tccatcattt ctcttctaag cttatcttcc aaaacatcca caagagcaag ttcattcagcc | 420 |
| agagactttc ccagaa | 436 |

<210> 1077
<211> 256
<212> DNA
<213> Homo sapien

| | |
|--|-----|
| <400> 1077 | |
| ctgaagatta ataggaaaca gtgaaaaagc aacgtcctgt gatcagtaac tttaaagaca | 60 |
| agcttggttc tctctttctg gcaactactga cattcccacc attctagctt ccgaattctg | 120 |
| gaaaaagaga agatgattaa caaaaataga gaatgtagaa acttctgggt ttgtgcctac | 180 |
| aggattggca ccagaccctc agtgcctca tgcctccatct acaaggcagc accctctcca | 240 |
| gaggcagcca gggagg | 256 |

<210> 1078

<211> 202
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(202)
 <223> n = A,T,C or G

<400> 1078
 ctgtgctncn caaccagatc catgtnaagt gccccgcccc gagaagggag ccaggggggag 60
 ctgactncag ncaacancca gtgnccggat gancaccaac atgtgagggg tgaaccttgg 120
 cctccangac atntgcaccc cctnccccacc tccaoggacc tgggacctcc aggcggctca 180
 gtgctgcctg cggcccagct aa 202

<210> 1079
 <211> 170
 <212> DNA
 <213> Homo sapien

<400> 1079
 gcgcttctcg ggcaccgtca ggcttaagtc cactccccgc cctaagttct ctgtgtgtgt 60
 cctgggggac cagcagcact gtgacgaggc taaggccgtg gatatcccc acatggacat 120
 cgaggcgctg aaaaaactca acaagaataa aaaactggtc aagaagctgg 170

<210> 1080
 <211> 494
 <212> DNA
 <213> Homo sapien

<400> 1080
 cctgcggcaa agagatgcgc ttattgagaa acatggctta gttataatcc ccgatggcac 60
 tcccaatggt gatgtcagtc atgaaccagt ggctggagcc atcactgttg tgtctcagga 120
 agctgctcag gtcttggagt cagcaggaga agggccatta gatgtaaggc tacgaaaact 180
 tgctggagag aaggaagaac tactgtcaca gattagaaaa ctgaagcttc agttagagga 240
 ggaacgacag aaatgctcca ggaatgatgg cacagtgggt gacctggcag gactgcagaa 300
 tggctcagac ttgcagttca tcgaaatgca gagagatgcc aatagacaaa ttagcgaata 360
 caaatttaag ctttcaaaag cagaacagga tataactacc ttggagcaaa gtattagccg 420
 gcttgagggg caggttctga gatataaaac tgctgctgag aatgctgagg aaagtgaag 480
 atgaattgaa agca 494

<210> 1081
 <211> 123
 <212> DNA
 <213> Homo sapien

<400> 1081
 ctgctgctat taagttgcaa gctctacagc tagctacatg actgatggat cagtttgaga 60
 tttgttcctt tgtcaaaaagt ttaactctga tagaaggttg gcctcacatt ctgatgtttg 120
 gac 123

<210> 1082
 <211> 297
 <212> DNA

<213> Homo sapien

<400> 1082

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctgcacttg | aacatggctt | tggttttaag | caacttctct | accctgaccc | tcctcctggg | 60 |
| acagcgtttc | gggaggtttc | ttggcctcac | tgagagggat | gtggagctgc | tgtaccccg | 120 |
| caaggagaag | gtattctaca | gcctgatgag | ggagagcggc | tacatgcaca | tccagtgac | 180 |
| caagcctgac | accgtaggct | ctgctctgaa | tgactctcct | gtgggtctgg | ctgcctatat | 240 |
| tctagagaag | ttttccacct | ggaccaatac | ggaattccga | tacctggagg | atggagg | 297 |

<210> 1083

<211> 452

<212> DNA

<213> Homo sapien

<400> 1083

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgggccacg | aggacaccac | cagcttggat | cggcctcgcc | gtgtggaata | ctttgtagat | 60 |
| aagcaactcc | aagtaaaggc | tgtcacctgt | gggcctggga | acacctacgt | gtatgctgtg | 120 |
| gagaaagga | agagctgaca | tgtgtacgta | tatgtatatg | caacacctgt | gagaccccca | 180 |
| ttcaggtcaa | ggaaaaccat | tgctgcacc | ccaaggggcc | catatttgcc | cctccccatc | 240 |
| acagtctgc | ccttcaccct | caagcacggt | cctaaacttg | tctgcacttt | agaaacacct | 300 |
| ggagagcatt | gaaaactctg | ctgcctaagg | tcagcatcaa | tcaaaacaat | gaaatcaatg | 360 |
| aaacaatgaa | accagagctt | ctagggtgtg | ggcctggata | gtggtagatt | caaagctcca | 420 |
| cccacctcat | cccaggtaca | tttgatgtgc | ag | | | 452 |

<210> 1084

<211> 301

<212> DNA

<213> Homo sapien

<400> 1084

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtagttga | ctgaagtcgc | taaacaggac | ggatttaagt | agaggtgata | tgtccagtca | 60 |
| cggcataga | gacgtcctct | gcgtcaccat | ccacacacag | ggcttctggt | agacatcg | 120 |
| caaagctctc | catgttaata | ttcatctgaa | tatggataat | taggggtggc | agcaaaacta | 180 |
| tcaactgtta | aatagtggag | atttctgtct | aggccatcta | tggctttcat | gtcctctgca | 240 |
| gtcaactgga | actcaaaaac | ctgcacgttc | tgtctgatgc | gctgctcatt | gtagctcttg | 300 |
| g | | | | | | 301 |

<210> 1085

<211> 369

<212> DNA

<213> Homo sapien

<400> 1085

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ctgtttccca | tgggccacca | ggcggctcag | gacagcaaac | gtctcatccc | ctctcaggat | 60 |
| gtacttctcc | atgtcctgct | cgatccactg | gtacatgagg | cccttcacat | gcacgtctcg | 120 |
| gatggcgctc | gtcacgtcct | tgtagagatg | tgcttgggtca | aactccaggc | tgtggcccag | 180 |
| aaagtagtcc | accacacagg | acagcagagc | catctccggt | agcgagaaga | tgtccatgaa | 240 |
| ctgcttaatg | gagggaccct | tgccatagaa | gccactcctc | tggtatagtg | ggatgtgctg | 300 |
| ggtaccccca | tacagctcaa | tcacctcctc | gtctggcaca | ggctggaggc | ccctgtaggc | 360 |
| tgtccccag | | | | | | 369 |

<210> 1086

<211> 316

<212> DNA

<213> Homo sapien

<400> 1086
 cctcagaggt ttctccacag tcctcttctg ggcaaattct tgtttcttca catgccggac 60
 tagcttaaga ccaatgcagt agcttatttc caagccttgc aaagtatata atatctaaga 120
 ggaaagggtt ttgtcatcca gcgttgtcca ctttgtgggg ctttgtaggt agacggagcc 180
 acactacagg caggggtatga gcagagggat gtatggagtg tgggtgactc tgagcctcac 240
 tgccgctgca aggtggggaa actgtaagtg aaccctgtg ggtgcggggg agggatatccg 300
 gtgcgcaggg aggtgg 316

<210> 1087

<211> 329

<212> DNA

<213> Homo sapien

<400> 1087
 cctgcagggg atgggacctt ccagaagtgg gcgtctgtgg tgggtgccttc tggacaggag 60
 cagagataca cctgccatgt gcagcatgag ggtctgcccc agccctcac cctgagatgg 120
 gacccgtctt cccagcccac catccccatc gtgggcatca ttgctggcct ggttctcttt 180
 ggagctgtga tcgctggagc tgtggtcgct gctgtgatgt ggaggaggaa gagctcagat 240
 agaaaaggag ggagctactc tcaggctgca agcagtgaca gtgcccaggg ctctgatatg 300
 tctccacag cttgtaaagt gtgagacag 329

<210> 1088

<211> 342

<212> DNA

<213> Homo sapien

<400> 1088
 ccactcactg ctgggaccca ggcacctccc ttctccatcc tctctggatt gtcagtaatg 60
 tcctggaaca gaagcctgtg ggatggcctt gggcacggag aagccctggg gtcagtgtcg 120
 tgcacggatg gcggcagtgt tgaaccacag aggctgaacc cggcccacca cggaagatga 180
 gtgcatggca accgcctgcc ttcacgtcgc tccacttggg aacccaagg tctgggctgt 240
 tctaggtatt gcttcacgtg ccccagcaag cccttaacaa gagggcctgg ttccctgaag 300
 aaccaatccc aggaaggggc cttgatccct ccgccttgct ga 342

<210> 1089

<211> 51

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(51)

<223> n = A,T,C or G

<400> 1089

ccttggtgtc agtctccncg ctcttcttgc cactgttgag ggtggagatg t 51

<210> 1090

<211> 515

<212> DNA

<213> Homo sapien

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(610)
<223> n = A,T,C or G

<400> 1094
ccatgcaaaa ggaggtggtg cactcagtgc agtcgctgcc acaaaaagtc cgattatattt 60
cattggtaca ggggaacata tagatgactt tgaacctttc aaaacacagc cttttatttag 120
caaacttctt ggtatgggcg acattgaagg actgatagat aaagtcaacg agttgaagtt 180
ggatgacaat gaagcactta tagagaagtt gaaacatggt cagtttacgt tgcgagacat 240
gtatgagcaa tttcaaaata tcatgaaaat gggccccttc agtcagatct tggggatgat 300
ccctggtttt gggacagatt ttatgagcaa aggaaatgaa caggagtcaa tggcaaggct 360
aaagaaatta atgacaataa tggatagtat gaatgatcaa gaactagaca gtacggatgg 420
tgccaaagtt tttagttaaac aaccaggaag aatccaaaga gtagcaagag gatcgggtgt 480
atcaacaaga gatgttcgag aacttttgac acaatatacc aagtttgac agatggtaaa 540
aaagatggga ggtatcaaag gacttttcaa aggtgggcga catgtctaan aatgtgagcc 600
agtcacagat 610

<210> 1095
<211> 232
<212> DNA
<213> Homo sapien

<400> 1095
ccttattttct cttgtccttt cgtacagga ggaatttgaa gtagatagaa accgacctgg 60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta 120
atagcggctg caccatcggg atgtcctgat ccaacatcga ggtcgtaaac cctattgttg 180
atatggactc tagaatagga ttgcgctggt atccctaggg taacttggtc cg 232

<210> 1096
<211> 377
<212> DNA
<213> Homo sapien

<400> 1096
ccacgctcat ggaaaccacc caaggacagc cagagtcac attccctggc aagctgggtg 60
tattcttcca aaagtttccc acccagtggt tcagacaggt gtagcgtctc tgcaggggtcc 120
cgtgcaatga agtcaaagtc ctcaggcagg aaagccaggc aggcacccag tctggcagcc 180
tctcgaacca gccacgcaca tgttttaaag ttctgttgct tgtctggcgt cgatgttacc 240
tggcacacag ccaccagggg cagttcgcag gaggaagagg agatagccat ggctctgggc 300
ctgggctgag cacaaggtac tgagagttga ggtatccgga gtccaggaca cagaaggagc 360
aggaatctgt gaggagg 377

<210> 1097
<211> 311
<212> DNA
<213> Homo sapien

<400> 1097
ccacgccatg gggctggagc actcccaaga ccctggggcc ctgatggcac ccatttacac 60
ctacaccaag aacttccgtc tgtcccagga tgacatcaag ggcattcagg agctctatgg 120

ggcctctcct gacattgacc ttggcaccgg cccaccccc acactggggc ctgtcactcc 180
 tgagatctgc aaacaggaca ttgtatttga tggcatcgct cagatccgtg gtgagatctt 240
 cttcttcaag gaccggttca tttggcggac tgtgacgcca cgtgacaagc ccatggggcc 300
 cctgctggtg g 311

<210> 1098

<211> 404

<212> DNA

<213> Homo sapien

<400> 1098

ccacccacgc ttaggttccc atcacactga tgactccggg tttggcgagc acaggagcgc 60
 aaaccttttc acattctttc tgtgatccaa atttgttttc gtttccacca caacctccat 120
 accagaatct tgcacagctt ttggtgtttg gatcatagta ccattttaat atgaaatccc 180
 tgcaagttcc ttcgtctttc ggcaacttgc atatatctgt ttcagtgaga gccaatgggt 240
 ctgtgctcac cattagattg atggttgaac tagaagctga ccttgctggc tgtggagggtg 300
 ggggctgaga tttctttgta ctgaaacttc cgtggtagggt ggctctgacc tgagacctca 360
 ggtagcagac cacagccaca tggatatgtct gcccagcgag cagg 404

<210> 1099

<211> 442

<212> DNA

<213> Homo sapien

<400> 1099

ccatgggatg gctcttctga ccattggggg ccaggccagg ccaggccagg cttagggtag 60
 caaggaccag gccaaagggg cagggcctcc tttggagggg ttgaggggta catcctcggc 120
 tgggtgtttgc atccaggggt ccagcaggat ctcttccagt gagggtcggg aagaaggttt 180
 gggggccagg caccggcgga ttagggcaca gcagtctggg gagacatggg ctgggaagtg 240
 gagctcagct tccagaatct cctggtccct ctcaaaggga atgtccccac acaccatgtc 300
 atagaggagg atgccagtg accagacagt ggccgggagt gcatgggtact ggtgtcgaga 360
 gatccactct ggggggctgt acacccttgt cccatcaaag tcagtgtagg gttcatcatg 420
 aagcagggca ccaggaacca aa 442

<210> 1100

<211> 191

<212> DNA

<213> Homo sapien

<400> 1100

ccacgaaaat caatgagaag ccacaggtga tcgcggacta tgagagcgga cgggccatac 60
 ccaataacca ggtgcttggc aaaatcgagc gggccattgg cctcaagctc cggggaaagg 120
 acattggaaa gcccatcgag aaggggccta gggcgaaatg aacacaaagc ctcgaaatca 180
 gtgcgctcca g 191

<210> 1101

<211> 178

<212> DNA

<213> Homo sapien

<400> 1101

cgggtacttt ggtggacatg aaggaactgg gcatatggga gccattggct gtgaagctgc 60
 agacttataa gacagcagtg gagacggcag ttctgctact gcgaattgat gacatcgttt 120
 caggccacaa aaagaaaggc gatgaccaga gccggcaagg cggggctcct gatgctgg 178

<400> 1105
ctggggccac tgtcggcacc atgattggag tgctgggttg ggttgctctg atatagcagc 60
cctgggtgtag tttcttcatt tcaggaagac tgacagttgt tttgcttctt ccttaaagca 120
tttgcaacag ctacagtcta aaattgcttc tttaccaagg atatttacgg aaaagactct 180
gaccagagat cgagaccatc ctagccaaca tcgtgaaacc ccatctctac taaaaatata 240
gaaattagct ggacatggtg gcatgtgcct gtaatccag ctactcagga ggctgaggca 300
ggagaactgc ttgaacaggg acccgggagg cggagatttg agtgagccga gatcgcgcca 360
ctgcactcca gtctgggcta cacagtgaga ctctgtctca agaaaaataa acagaagaat 420
tgggggttg ggggtgggaaa cagtgtttcc aggcagagag aacagcacgt acaaaggaga 480
ctgttgggag gggttaaatga aataattcat gtaagggtact tagtaccaca catgaatttc 540
acaagcagca g 551

<210> 1106
<211> 280
<212> DNA
<213> Homo sapien

<400> 1106
ctgctcttca cacagggttc tggggaaaaac aaggaagaga tcatcaatta tgaatttgac 60
accaaggacc tgggtgtgcct gggcctgagc agcatcggtg gcgtctggtta cctgctgagg 120
aagcactgga ttgccaacaa cctttttggc ctggccttct cccttaatgg agtagggctc 180
ctgcacctca acaatgtcag cactggctgc atcctgctgg gcggactctt catctacgat 240
gtcttctggg tatttggcac caatgtgatg gtgacagtgg 280

<210> 1107
<211> 570
<212> DNA
<213> Homo sapien

<400> 1107
ctgattagtg tctaaggaat ggtccaatac tgttgccctt ttccttgact attacactgc 60
ctggaggata gcagagaagc ctgtctgtac ttcattcaaa aagccaaaat agagagtata 120
cagtcctaga gaattcctct atttgttcag atctcataga tgacccccag gtattgtctt 180
ttgacatcca gcagtccaag gtattgagac atattactgg aagtaagaaa tattactata 240
attgagaact acagctttta agattgtact tttatcttaa aagggtggtta gttttcccta 300
aaatacttat tatgtaaggg tcattagaca aatgtcttga agtagacatg gaatttatga 360
atggttcttt atcattttct tttccctttt ttggcatcct ggcttgccct cagttttagg 420
tccttttagt tgcttctgta agcaacggga acacctgctg agggggctct ttcctcatg 480
tatacttcaa gtaagatcaa gaatcttttg tgaaattata gaaatttact atgtaaatgc 540
ttgatggaat tttttcctgc tagtgtagct 570

<210> 1108
<211> 386
<212> DNA
<213> Homo sapien

<400> 1108
ctgttctctg ggtgacactg tataaacacg atgaccctgc cttgacttta gttgctggtc 60
ttacatcaaa taagcccaca gacaaactcc gtgccctgcc tctgtggtta tctttacaat 120
acttgggact tgatggggtt gtggagagga tcaagcatgc ctgtcaactg agtcaacggg 180
tgcaggaaaag tttgaagaaa gtgaattaca tcaaaatctt ggtggaagat gagctcagct 240
ccccagtgg ggtgttcaga tttttccagg aattaccagg ctgagatccg gtgttttaaag 300
ccgtcccagt gcccacatg acaccttcag gagtcggccg ggagaggcac tcgtgtgacg 360
cgctgaatcg ctggctggga gaacag 386

<210> 1109
 <211> 409
 <212> DNA
 <213> Homo sapien

<400> 1109
 ctctggtctg taaccagtct cttcaaggca ttatctcctg gggccaggat ccgtgtgcga 60
 tcacccgaaa gcctggtgtc tacacgaaag tctgcaaata tgtggactgg atccaggaga 120
 cgatgaagaa caattagact ggacccaccc accacagccc atcacccctcc atttccactt 180
 ggtgttttgt tcctgttcac tctgttaata agaaacccta agccaagacc ctctacgaac 240
 attctttggg cctcctggac tacaggagat gctgtcactt aataatcaac ctgggggttcg 300
 aaatcagtga gacctggatt caaattctgc cttgaaatat tgtgactctg ggaatgacaa 360
 cacctgggtt gttctctgtt gtatccccag ccccaaagac agctcctgg 409

<210> 1110
 <211> 215
 <212> DNA
 <213> Homo sapien

<400> 1110
 ccattttgga gtgtgtccat tgggtagcaa tgtggaaacc accagggcct ttgtggagaa 60
 aatggagggg gttgaggag tcccaggagg ggcttatttg agggcctttg ccacttgctc 120
 ataggcgagc tcgatctcct catcatctgg acaggtggaa gcgaattctt cccgggcgta 180
 ggcattgctc aagtaccgat gcactccccg gaagg 215

<210> 1111
 <211> 308
 <212> DNA
 <213> Homo sapien

<400> 1111
 cctgggcccg ctgacttcag ggtgaggcca cagctactgc agcgcttttt atttatttat 60
 ttatttactg agatggagtc ttgctctgtc acccaggctg gagtgcagtg gtgcaatctc 120
 ggctcactgc aacctctgcc tcctgggctg cagtgattct cctgcgttca agtaattctc 180
 ctgcctcggc cttctgagta gttgggatta caggcatatg ccaccacact tggctaattt 240
 tttgtatttt tagtagaaat ggggtttcac catgttggcg aggctggtct cgaactcctg 300
 acctcaag 308

<210> 1112
 <211> 177
 <212> DNA
 <213> Homo sapien

<400> 1112
 ccactggctc cctgggccag ggccctgggg cgccttgtgg gatggcctac accggcaaatt 60
 acttcgacaa ggccagctac cgagctctatt gcttgctggg agacggggag ctgtcagagg 120
 gctctgtatg ggaggccatg gccttcgcca gcctctataa gctggacaac cttgtgg 177

<210> 1113
 <211> 646
 <212> DNA
 <213> Homo sapien


```
<400> 1123
ccaattgaaa caaacagttc tgagaccgtt cttccactac tgattaagag tgggggtggca      60
```

```
<210> 1124
<211> 219
<212> DNA
<213> Homo sapien
```

```
<210> 1125
<211> 246
<212> DNA
<213> Homo sapien
```

```
<210> 1126
<211> 227
<212> DNA
<213> Homo sapien
```

```
<210> 1127
<211> 377
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|------------|------------|------------|-------------|--|-----|
| <400> | 1127 | | | | | | |
| cttgccgtcg | atgccaggga | ggcgcacagg | accttctttt | ccagcggggc | cgatatatttc | | 60 |
| aggggaacca | ggaagacctc | tgggtcccat | gagaccaggc | tccccagggc | gaccagcatc | | 120 |
| tccattaggt | cctcggactc | cagcagggcc | acttgacca | cgactaccag | gagggcccat | | 180 |
| gacgccagct | ctgccatcag | ctccaggaag | accacgagaa | ccaggactac | ctctcagccc | | 240 |
| aggaggtcct | ggagggccgg | cagatccagc | ttccccatta | gggcctctct | ttccttcttc | | 300 |
| accactggga | ccaggaggac | cttggggccc | agcagagccg | ggctcaccct | tgttaccgct | | 360 |

377

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<211> 304

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<400> 1136
cctgccgtcg atgccagga ggccgacagg accttctttt ccagcggggc cgatatattcc 60

```

aggggaacca ggaagacctc tgggtcccat gagaccaggc tccccagggc gaccagcatc 120
tccattaggt cctcggactc cagcagggcc acttgacca cgactaccag gagggcccat 180
gacgccagct ctgccatcag ctccaggaag accacgagaa ccaggactac ctctcagccc 240
aggaggtcct ggagggccgg cagatccagc tccccatta gggectctct ttccttcttc 300
accactggga ccaggaggac cttggggccc agcagagccg ggctcaccct tgttaccgct 360
ctctcctttg gagccag 377

```

<210> 1137

<211> 250

<212> DNA

<213> Homo sapien

<400> 1137

```

ctgttcaact tccaactcta aataggcacc attaaacaaa aaaccccagt attttaaatt 60
tctccagcac acattccagg atcaatgctc tgaactgtaa tcagctagta attcataacg 120
ggaatacagc cttagaatgg aagctatatt gcttccctgc cccctttctc ttacaattgg 180
agagtgtagg tattaaggga taaaaagtca gaggaagaat aattaaaaag aaaaatgccc 240
aaagctgcag 250

```

<210> 1138

<211> 511

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(511)

<223> n = A,T,C or G

<400> 1138

```

tcgaccaggt cctcctgggc catctggtcc ccgagggtcag cctgggtgtca tgggcttccc 60
cggctcctaaa ggaaatgatg gtgctcctgg taagaatgga gaacgagggtg gccctggagg 120
acctggccct cagggtcctc ctggaaagaa tggtgaaact ggacctcagg gacccccagg 180
gctactggg cctgggtggtg acaaaggaga cacaggaccc cctgggtccac aaggattaca 240
aggcttgctt ggtacagggtg gtccctcagg agaaaatgga aaacctgggg aaccagggtc 300
aaaggggtgat gccgggtgcac ctggagctcc aggaggcaag ggtgatgctg gtgcccctgg 360
tgaacgtgga cctcctggat tggcaggggc cccaggactt agagggtggag ctgggtcccc 420
tgggtcccga ngaggaaagg gtgctgctgg tcctcctggg ccacctggtg ctgctggtac 480
tcctggtctg caaggaatgc ctggagaaag a 511

```

<210> 1139

<211> 505

<212> DNA

<213> Homo sapien

<400> 1139

```

ctgtggactc cagcatgttt ctgataatta tgcaagcaac aattctgtag cctcaagtaa 60
gaccacctgt gaacttgatc attatctggc ccaaatatga agataaaacta taactttgga 120
gtttgtttcc tatttgtatt cacattctgc ttctaaatc agttttctaa attgtgcttg 180
caattaggca ttggtcaggg gtgaatggct cttttcacag agagtagcca accagagacc 240
tttgctttga tatcatcaac tgcagagaat gctgttgatg ggaatgctgg aagcagaaac 300
tttgtcatcg gaaaaacttt tcttgtatgc atgagactca acatcaggat ccacagctta 360
aagatgggaa ttcagggtatg aaagaaaaca ggcaaggagg cactgaggga gaaagacaca 420
gacttttatc ctctgtggct cattgttact ggaatattct aaaactcttg ttcacatgct 480

```

attatgactt ataaagcagc aacag

505

<210> 1140

<211> 256

<212> DNA

<213> Homo sapien

<400> 1140

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtagcttc | tgtgggactt | ccactgctcg | ggcgtcaggc | tcaggtagct | gctggccgcg | 60 |
| tacttggtgt | tgctctgttt | ggagggtttg | gtggtctcca | ctcccgctt | gacggggctg | 120 |
| ccatctgcct | tccaggccac | tgtcacagct | cccggttaga | agtcactgat | cagacacact | 180 |
| agtgtggcct | tgttggcttg | gagctcctca | gaggagggcg | ggaacagagt | gacagtgggg | 240 |
| ttggccttgg | gctgac | | | | | 256 |

<210> 1141

<211> 371

<212> DNA

<213> Homo sapien

<400> 1141

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ccaggggccc | attctgtctg | tgggactgtg | ggttctcagt | ggaattgttg | cctttcttgt | 60 |
| cgtggagaaa | tttgtgagac | atgtgaaagg | aggacatgg | cacagtcag | gacatggaca | 120 |
| cgctcacagt | catgcacgtg | gaagtcacgt | acatggaaga | caagagcgtt | ctaccaagga | 180 |
| gaagcagagc | tcagaggaag | aagaaaagga | aacaagaggg | gttcagaaga | ggcgaggagg | 240 |
| gagcacagta | cccaaagatg | ggccagttag | acctcagaac | gctgaagaag | aaaaaagagg | 300 |
| cttagacctg | cgtgtgtcgg | ggtacctgaa | tctggctgct | gacttggcac | acaacttcac | 360 |
| tgatgggtctg | g | | | | | 371 |

<210> 1142

<211> 312

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(312)

<223> n = A,T,C or G

<400> 1142

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctccacac | tgtcaaagt | caactccacc | agcactgaga | caatgagtag | atgagaatgt | 60 |
| agaaagaggg | aaggtggtag | gtaaaggagc | ggaaggaaga | ggtggggaaa | gaggggaagg | 120 |
| ggtaggtaaa | ggagcggaag | gaagaggtgg | ggaaagaggg | aaggagagaa | gggaaggagg | 180 |
| gaagagaaa | aaggaagaaa | aggaaagcat | ggcccggtta | gagacaaagc | cagaggtgat | 240 |
| caggtcagca | gcaggagagg | ctcagaaggg | agcctctcgg | gaagtgcagg | cngccatgag | 300 |
| ggctcgtttc | ag | | | | | 312 |

<210> 1143

<211> 367

<212> DNA

<213> Homo sapien

<400> 1143

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccagacgtgg | tggctcacac | ctgcaatccc | agcaccttag | gaggccgagg | caggaggatc | 60 |
| cttgaggtca | ggagttcgag | accagcctcg | ccaacatggt | gaaaccccat | ttctactaaa | 120 |

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| atacaaaaaa | ttagccaagt | gtggtggcat | atgcctgtaa | tcccaactac | tcagaaggcc | 180 |
| gaggcaggag | aattacttga | acgcaggaga | atcactgcag | cccaggaggc | agaggttgca | 240 |
| gtgagccgag | attgcaccac | tgcactccag | cctgggtgac | tgagcaagac | tccatctcag | 300 |
| taaataaata | aataaataaa | aagcgctgca | gtagctgtgg | cctcaccctg | aagtcagcgg | 360 |
| gcccagg | | | | | | 367 |

<210> 1144

<211> 159

<212> DNA

<213> Homo sapien

<400> 1144

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctggaggag | cggccgcaca | cacagccagg | cgctaggctc | cctgcgggac | ctcgggaagg | 60 |
| gggaagagcg | tcaacgattt | acggagggtc | cagccgctgg | gtcagattga | gacaaaccat | 120 |
| tgtgtggttg | ggttcgggtc | agcaggctgg | agagggttc | | | 159 |

<210> 1145

<211> 450

<212> DNA

<213> Homo sapien

<400> 1145

| | | | | | | |
|------------|------------|-------------|------------|------------|-------------|-----|
| ccatgggtgt | ctggagcacc | ctgaaactgt | atcaaagttg | tacatatttc | caaacatttt | 60 |
| taaaatgaaa | aggcactctc | gtgttctcct | cactctgtgc | actttgctgt | tgggtgtgaca | 120 |
| aggcatttaa | agatgtttct | ggcattttct | ttttatttgt | aagggtggtg | taactatggt | 180 |
| tattggctag | aaatcctgag | ttttcaactg | tatatatcta | tagtttgtaa | aaagaacaaa | 240 |
| acaaccgaga | caaacccttg | atgctccttg | ctcggcgttg | aggctgtggg | gaagatgcct | 300 |
| tttgggagag | gctgtagctc | agggcggtgc | ctgtgaggct | ggacctgttg | actctgcagg | 360 |
| gggcatccat | ttagcttcag | gttgtcttgt | ttctgtatat | agtgacatag | cattctgctg | 420 |
| ccatcttagc | tgtggacaaa | gggggggtcag | | | | 450 |

<210> 1146

<211> 324

<212> DNA

<213> Homo sapien

<400> 1146

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ccatacaggg | ctgttgccca | ggccctagag | gtcattcctc | gtaccctgat | ccagaactgt | 60 |
| ggggccagca | ccatccgtct | acttacctcc | cttcggggcca | agcacaccca | ggagaactgt | 120 |
| gagacctggg | gtgtaaatgg | tgagacgggt | actttggtgg | acatgaagga | actgggcata | 180 |
| tgggagccat | tggctgtgaa | gctgcagact | tataagacag | cagtggagac | ggcagttctg | 240 |
| ctactgcgaa | ttgatgacat | cgtttcaggc | cacaaaaaga | aaggcgatga | ccagagccgg | 300 |
| caaggcgggg | ctcctgatgc | tgga | | | | 324 |

<210> 1147

<211> 191

<212> DNA

<213> Homo sapien

<400> 1147

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccacgaaaat | caatgagaag | ccacaggtga | tcgcggacta | tgagagcgga | cggggccatac | 60 |
| ccaataacca | ggtgcttggc | aaaatcgagc | gggccattgg | cctcaagctc | cggggaaagg | 120 |
| acattggaaa | gcccacgag | aaggggccta | gggcgaaatg | aacacaaagc | ctcgaaatca | 180 |
| gtgtgctcca | g | | | | | 191 |

<210> 1148
 <211> 344
 <212> DNA
 <213> Homo sapien

<400> 1148
 ctgtccaatg acaacaggac cctcactota ctcagtgtca caaggaatga tgtaggaccc 60
 tatgagtgtg gaatccagaa cgaattaagt gttgaccaca gcgaccaggt catcctgaat 120
 gtccctctatg gccagacga ccccaccatt tccccctcat acacctatta ccgtccaggg 180
 gtgaacctca gcctctcctg ccatgcagcc tctaaccacac ctgcacagta ttcttggtctg 240
 attgatggga acatccagca acacacacaa gagctcttta tctccaacat cactgagaag 300
 aacagcggac tctatacctg ccaggccaat aactcagcca gtgg 344

<210> 1149
 <211> 329
 <212> DNA
 <213> Homo sapien

<400> 1149
 ctgaccact cactgggcgg gggcacaggc tctggaatgg gcactctcct tatcagcaag 60
 atccgagaag aataccctga tcgcatcatg aataccttca gtgtggtgcc ttcacccaaa 120
 gtgtctgaca ccgtggtcga gccctacaat gccaccctct ccgtccatca gttggtagag 180
 aatactgatg agacctattg cattgacaac gaggccctct atgatattctg ctccgcact 240
 ctgaagctga ccacaccaac ctacggggat ctgaaccacc ttgtctcagc caccatgagt 300
 ggtgtcacca cctgcctcgg tttccctgg 329

<210> 1150
 <211> 406
 <212> DNA
 <213> Homo sapien

<400> 1150
 ccagttatatt gcaagtggta agagcctatt taccataaat aatactaaga accaactcaa 60
 gtcaaaccctt aatgccattg ttattgtgaa ttaggattaa gtagtaattt tcagaattca 120
 cattaacttg attttaaaat cagttttgtg agtcatttac cacaagctaa atgtgtacac 180
 tatgataaaa acaaccattg tatcctgtt tttctaaaca gtactaattt ctaacactgt 240
 atatatcctt cgacatcaat gaactttgtt ttcttttact ccagtaataa agtaggcaca 300
 gatctgtcca caacaaactt gccctctcat gccttgctc tcaccatgct ctgctccagg 360
 tcagccccct tttggcctgt ttgttttgtc aaaaacctaa tctgct 406

<210> 1151
 <211> 346
 <212> DNA
 <213> Homo sapien

<400> 1151
 ctgcgtgagt accaggagct gatgaacgtc aagctggccc tggacatcga gatcgccacc 60
 tacaggaagc tgctggaggg cgaggagagc cggctggagt ctgggatgca gaacatgagt 120
 attcatatga agaccaccag cggctatgca ggtggtctga gctcggccta tgggggacctc 180
 acaagccccg gcctcagcta cagcctgggc tccagctttg gctctggcgc gggtccagc 240
 tccttcagcc gcaccagctc ctccagggcc gtggttgtga agaagatcga gacacgtgat 300
 gggaagctgg tgtctgagtc ctctgacgtc ctgcccgaagt gaacag 346

<210> 1152
 <211> 427
 <212> DNA
 <213> Homo sapien

<400> 1152
 ctggactgct gtacatcaag gacagattaa ctggaaaaca tatgttcctt atgcgtgac 60
 gagagccatt cagaaaagac ttcctttgtg ttcagcctat acttttccat atgggtatacc 120
 ttgaaaaaaa ttagcacacc atgggttattt ttctaccttt tataaaagac agagcctgtt 180
 tactcattta gaagatagag aaaattgggtc taaaattgaa catcctagat tcacactccc 240
 aagtcactta aggtgatttg atggtgagga aaatgattga cagagcccaa caatgatctc 300
 aggaattaca ttttccaaca gaccaaaaaa tgttttcatg tagcagcaat gcagatttgg 360
 tgaatattta atatataatt tagtatgtat ttcactttat gactgacaat taaaaaatat 420
 tgtttgg 427

<210> 1153
 <211> 331
 <212> DNA
 <213> Homo sapien

<400> 1153
 ctggccggcg gtgcagatct ggagtccagc ctcagggatg cgctactttc cattctctgc 60
 attgaacatt cgttctgtca gcatccgctc cagcttcaact gcatcagcgg caaacttgcg 120
 gatcccgta gagagcttct ccacagccat ctggctcctcg ttgtgcaacc aacggaaaga 180
 cttctcatcc aggtggattt tttccaggtc actggcttgg gctggggggac aagaaccagc 240
 cttccatgcc tgctccatgt ccctgcccac cttggccctt tgggctcagg gctgaaccg 300
 ctgcacccaa gcatctccca ccagggccag g 331

<210> 1154
 <211> 403
 <212> DNA
 <213> Homo sapien

<400> 1154
 ctgaactttc agatgaagtt gacttctact tgattgcagg attcaggggtt tctcagatgt 60
 taatacagag tcaaaagcgg tggataaaac cttgcaaatt gcttgtgctt gttccaggct 120
 gttgactga taaaccaca ggctgtattc ctcattgctt gcatctgtgg tcttcagagc 180
 cagtaagctt tttcccgccc ccagaccgtc atcgtaacac accatccgga ttattaagta 240
 gagagcatgc ctgtgcaaaa catcatattg atctgatgtt gatactttta tgccatactt 300
 ggaaactccc ataataaatt cttcctccgg aggaacaaaa ggcaactttc catcttgctg 360
 ggcaacgtct atataattta tcagggtctaa tggcccttca agg 403

<210> 1155
 <211> 491
 <212> DNA
 <213> Homo sapien

<400> 1155
 cctccctctc agagcttgcc ccagggactc tctggccctc agggttcaat gtattctgac 60
 caaggccaag ctttcctggg gctcaggga aatcacactt tgctacccga agctgtatcc 120
 cctcagatgc caggaaggcc gtgatcatct gactccacc ccttgagaca cattctctcc 180
 ctgactgtcc tgttctaagt cagcggagca ccttaggatg gaggggtgga ggcgaggcca 240
 gatgcagcct ctgtgaacag gtgcctggag gctgggaaat gaccctgaga ggcgaggaca 300
 cagcaaccgt gggcttaagg tgacctgag agcaagcttg gccacttta caattctgtt 360

```
<210> 1156
<211> 586
<212> DNA
<213> Homo sapien
```

```
<210> 1157
<211> 392
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(392)
<223> n = A,T,C or G
```

```
<210> 1158
<211> 375
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|------------|------------|------------|------------|--|-----|
| <400> | 1158 | | | | | | |
| gggaaaaata | attttatttc | tcaaatgata | agcacattca | gaagcaggac | agaggagctc | | 60 |
| tgatgacata | tctgggggac | tcaaaagggc | cctcattttc | tggtattttc | ccaggtgatt | | 120 |
| ctcttccaac | ctgtgagtc | tgtctctttt | cctcccatct | gaagtttgag | acatcctctg | | 180 |
| ccacaaggaa | agccaccaat | accagcccaa | agagccacca | gagaggaacc | aaaccacatg | | 240 |
| catcaagtta | taggaaggat | gcaagaaggg | aaattaggaa | ggaaagggag | gagtttagtt | | 300 |
| ggcattctgg | ggcatgctaa | catgagggcg | atggtctctc | tccaagtcgc | tggacataac | | 360 |
| ccttttcttt | ccagg | | | | | | 375 |

<210> 1163
<211> 337

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(337)
<223> n = A,T,C or G

<400> 1163
ctgcagagtg ggganaggct tttgccacta gaaacttcca ggatgcacga gatcaaggaa 60
ttaagtctgt aacaaaataa caggatgctc tgtgaagtcc aaagaattgc ttgaggcaaa 120
ctgcagagct ccatgagatc agcaacccca agagctttta caccgccgga cacggtttaa 180
taggaaaaaa atctcctata ctgnntattc anaaccaa at gaanagaaat gtcaaaggag 240
tcggaaacaa tatgtcaa at tangtaaatt cctgacctga cccanatttt gcngaacatt 300
tgatcctaaa ctgtgctgtc cacgtcctta ggatcac 337

<210> 1164
<211> 368
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(368)
<223> n = A,T,C or G

<400> 1164
ccagacgtgg tggctcacac ctgcaatccc agcaccttag gaggccgagg caggaggatc 60
cttgagggtca ggagttcgag accagcctcg ccaacatggg gaaaccccat ttctactaaa 120
aatacaaaaa attagccaag tgtgggtggca tatgcctgta atcccaacta ctccagaaggc 180
cgaggcgagga gaattacttg aacgcaggag aatcactgca ncccangagg canagggttg 240
antgagccga gattgcacca ctgcactcca gcctgggtga cagagcaaga ctccatctca 300
gtaaataaat aaataaataa aaagcgctgc agtagctgtg gcctcaccct gaagtcagcg 360
ggccccagg 368

<210> 1165
<211> 267
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(267)
<223> n = A,T,C or G

<400> 1165
ctgggaagga ggctcctccg ctttctcctg tttgtcatcc tctcatcag actcgacctc 60
catctcaact tctcactct ccccaaactt ttcatagcgc tctgaatga ggattcgggc 120
cccagctcc tctggcgtgg tggggggagg gaagttccct tgctcattgg gttggaagnc 180
cactgtttcc accaccacaa aatcatgcc a ntcnatctga gcataggcca cccgntcctt 240
ctccttctcc nnttcttct tcttctt 267

<210> 1166
<211> 433

<212> DNA
<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(433)

<223> n = A,T,C or G

<400> 1166

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtctgtac | actttttctt | gggggaagag | ttcttgtctt | cagtttactg | cagtaggggt | 60 |
| cctggctctg | ttacatgctc | atgtgttccg | gaagaacaca | tgaaatatca | tcccacggat | 120 |
| gacgatacag | cccctgcttc | ancctcttct | gatcaagata | gtgtccaatg | aaccccatac | 180 |
| tccttcccag | cacaaagatg | ccattgaggg | ctccaatgtc | aatatatcca | tcagcttcc | 240 |
| ccctgcaaca | cacatcaact | tgtagtttta | aaagggtcac | gtgactgcc | tcctccccac | 300 |
| agacagtact | actactgcc | aanaatgaga | agaaaagggg | tgtctgtggg | ggtngcatta | 360 |
| caggcaattt | ttgttntctt | nnttatacct | ctccttattt | tncaaanttt | ctattatgag | 420 |
| tntgcattac | ttt | | | | | 433 |

<210> 1167

<211> 362

<212> DNA

<213> Homo sapien

<400> 1167

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctctggctc | tttcttcagc | caettctcca | gtcctgcag | gttctgggtc | gagtagtcag | 60 |
| tgacgacgat | ctccttaaag | gattcacaa | cagagaggag | ctgatagata | gtggggccag | 120 |
| agccgatgtc | aatcagcagg | tctcccttca | caccgtctag | gcagaatata | ttgaaaagat | 180 |
| ttttcagaag | gtgcttaaga | atctggcttt | ctgcagagtg | cctagaacca | aacttgtaat | 240 |
| atTTTTctag | gtaatcccga | gggttaaaat | ggcttagata | ggtgtccttg | gaggtgaagc | 300 |
| ctgattccat | tatgtctcac | ttccgtacca | ctggagcact | gccctccttc | tctttcctcc | 360 |
| ag | | | | | | 362 |

<210> 1168

<211> 459

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(459)

<223> n = A,T,C or G

<400> 1168

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| gcagtcattg | ggcccaggac | catgccactg | gccctgctcc | cccagccgca | gcctcacctg | 60 |
| caggtgctcc | tcgatgtcct | tgcggtcgta | ggtgatgcca | ctgggcgtga | tgcaaggctc | 120 |
| ccgcatcagc | tcaaagctga | tcttgccaca | caggtagtcg | gggatgtctc | gcttctgtgg | 180 |
| cacaggggca | cacggtcaga | ggctgaaaag | gggcactgca | cgagcacctg | ccagccatcg | 240 |
| gcagcaagcg | acacacactc | accttctctt | tctcatccac | ctgagaaaaa | agctcgtcca | 300 |
| tgtccgccat | gtacttgtcc | tgtgaagagt | tgagtgtgtg | gcttggggga | gacacccac | 360 |
| ctccctcctn | catggggcac | anacccaaca | caaggcgggg | atgctnccac | gccacgtgca | 420 |
| cacacacaga | cccacatgtg | ggtggggggc | accctcacg | | | 459 |

<210> 1169

<211> 386

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaggccacc | tgtgcggggc | tctctgatgt | ggaaggttcg | ggtgaggaga | ttgtagaagg | 60 |
| agccgtagca | cacggccacc | acagtgcacg | tgaggcagat | cacgctgtag | ggcatgctga | 120 |
| agtccggtgt | cggcaggttc | accagcagcg | gctccgtgta | gagccgcaca | aagtagttag | 180 |
| agccatcaga | gactgggaac | aggctgttga | agaggggact | ctcttcccag | tccactggct | 240 |
| tggttgctac | catgctgggc | acaagggcgc | tgaggacaga | tgggctgaca | tagaagccat | 300 |
| ggttaggatc | tggcgtgtac | tcggtccact | tcagcagcgc | ccgctcaaac | tggatggaaa | 360 |
| ccttggtgac | tgagttggcc | ggccag | | | | 386 |

<213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|-------------|-------------|-----|
| ctatttctct | gttagtgttt | aaccaaccat | ctgttctaaa | agaagggtctg | aactgatgga | 60 |
| aggaatgctg | ttagcctgag | actcaggaag | acaacttctg | cagggtcact | ccctggcttc | 120 |
| tggaggaaaag | agaaggaggg | cagtgtctca | gtggtacaga | agtgagacat | aatggaatca | 180 |
| ggcttcacct | ccaaggacac | ctatctaagc | cattttaacc | ctcgggatta | cctagaaaaa | 240 |
| tattacaagt | ttggttctag | gcactctgca | gaaagccaga | ttcttaagca | ccttctgaaa | 300 |
| aatcttttca | agatattctg | cctagacggg | gtgaaggagg | acctgctgat | tgacatcggc | 360 |
| tctggcccca | ctatctatca | gctcctctct | gcttgatga | cctttaagga | gatcgctcgtc | 420 |
| actgactact | caggaccaga | acctgcagga | gctggagaag | tggctgaaga | aagagccaga | 480 |

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| cctcagcagc | cctgccacgg | atctgccga | ttctttcgca | tcaagaagtt | gatcttgcca | 60 |
| gccatttcca | tgttgtagat | ccgcgggcac | ctttcatagc | tttccctctg | tcgccggcgg | 120 |
| catggcttct | cataataccg | ccgatgctta | atgtcctcaa | tgagcccata | catagtgagg | 180 |
| attctgttta | gggtcctgta | tgcgctttcc | acgttccctt | cctgtaccat | cacagtctctg | 240 |
| gcgatgaact | tcagatgttt | tgccatgacc | ttggatttaa | accttcactc | tgtagagcct | 300 |
| cgcgcgctca | gtaccta | | | | | 317 |

<213> Homo sapien

<223> n = A, T, C or G

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| ggcaacgggga | ggaacagcag | cagaggcagc | angagcagga | ggagcgtgaa | cgagaagagc | 60 |
| ancggcgatn | ngctgcncctc | agtgaccgan | agaagagagc | tctggctgca | nagcgccgac | 120 |
| tcqctqccca | qttgggagcc | cctacctctc | caatccctga | ctctgcaatc | gtcaatactc | 180 |

gacgctgctg gagttgtggg gc

202

<210> 1173

<211> 173

<212> DNA

<213> Homo sapien

<400> 1173

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ctgcctgggt | tgtggccgcc | ctagcatcct | gtatgccac | agctactgga | atccccgctg | 60 |
| ctgctccagg | ccaagcttct | ggttgattaa | tgagggcatg | gggtgggtccc | tcaagacctt | 120 |
| cccctacctt | ttgtggaacc | agtgatgcct | caaagacagt | gtcccctcca | cag | 173 |

<210> 1174

<211> 301

<212> DNA

<213> Homo sapien

<400> 1174

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| ccaagagcta | caatgggcag | cgcacacagac | agaacgtgca | ggtttttgag | ttccagttga | 60 |
| ctgcgaggga | catgaaagcc | atagatggcc | tagacagaaa | tctccactat | tttaacagtg | 120 |
| atagttttgc | tagccaccct | aattatccat | attcagatga | atattaacat | ggagagcttt | 180 |
| gcctgatgtc | taccagaagc | cctgtgtgtg | gatggtgacg | cagaggacgt | ctctatgccg | 240 |
| gtgactggac | atatcacctc | tacttaaata | cgctcctgtt | agcgacttca | gtcaactaca | 300 |
| g | | | | | | 301 |

<210> 1175

<211> 537

<212> DNA

<213> Homo sapien

<400> 1175

| | | | | | | |
|------------|-------------|-------------|------------|-------------|-------------|-----|
| cctgcagggc | tggcccgtag | gagaagggtca | gggcccaggg | cttcagcagg | gggcacttgt | 60 |
| taatggcatt | gaggttgatg | gacgcctcct | cctcactctg | gcctccagac | aggaagggtga | 120 |
| tcccagtgac | agcggggggc | actgtgcggc | gcagcgctgt | gacgggtcgcc | atggcaatct | 180 |
| cctcatgaga | aaacttctga | gtgcaagcat | ggcctggggg | gacctggttg | ggcttcagca | 240 |
| aggtgccttc | caggtagatg | tggtggtcac | tcagagcctt | gtagacagca | gccagcacct | 300 |
| tctcggtcac | atactggcag | cgcttcaagt | catgggtccc | atcagggagg | atctcaggct | 360 |
| ccacgatggg | cacaatgccca | ttctgctggc | agatactggc | ataacggggc | agaacattgg | 420 |
| cattttccat | gatggcgagg | gctgaggggg | tgtgttcccc | aatcttcagc | acacaacgcc | 480 |
| acttggcgaa | gtcagctccg | tccttcttgt | actgggcaca | gcgctcagac | agcccat | 537 |

<210> 1176

<211> 384

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(384)

<223> n = A,T,C or G

<400> 1176

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ctgacaaaaa | atgtgaaatt | tccacaaaaa | atccaaactta | tgtgactaaa | cgcagtagtt | 60 |
| tttttaaaag | gggagataga | aaataaatgg | ttttgttgga | gtgcatttta | gtaagccttt | 120 |

```

gcagtaaaat gacggttgta actactaaac caaatttagt tttcacagca tggttttgtt 180
gttttcccct tgtttttcag aggtaaaattt tgcattatat ccttcagtat tttaacacta 240
ttttggcagt ttacacatta ctttttgntt ttccttcctt tttgngaaat gtattaagtt 300
gtggttctta ttgaaacagt attatataat gttngcttaa ttatatcatg tgatgctcan 360
ntctattntg atttattcat tagt 384

```

<210> 1177

<211> 562

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(562)

<223> n = A,T,C or G

<400> 1177

```

ccaacaacat gcaggaagct cagagtatcg atgaaatcta caaatacgac aagaaacagc 60
agcaagaaat cctggcgggc aagccctggg ctaaggatca ccattacttt aagtactgca 120
aaatctcagc attggctctg ctgaagatgg tgatgcatgc cagatcgagg ggcaacttgg 180
aagtgatggg tctgatgcta ggaaagggtg atggtgaaac catgatcatt atggacagtt 240
ttgctttgcc tgtggagggc actgaaaccc gagtaaatgc tcaggctgct gcatatgaat 300
acatggctgc atacatagaa aatgcaaaac aggttggccg ccttgaaaat gcaatcgggt 360
ggatcatag ccaccctggc tatggctgct ggctttctgg gattgatgtt agtactcaga 420
tgctcaatca gcagttccag gaaccatttg tagcagtggg gattgatcca acaagaacaa 480
tatccgcagg gnaaagtga tcttggcgcc tttaggacat acccaaaggg ctacaaacct 540
nctgatgaan gaccttctga gt 562

```

<210> 1178

<211> 353

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(353)

<223> n = A,T,C or G

<400> 1178

```

cgcgtctgga tggccgaatc attcgcacag actgggacgc aggctttaag gagggcaggc 60
aatacggccg tgggcatctt gggggccagg ttcgggatga gtatcggcag gactacnatg 120
ctgggagagg aggctatgga aaactggcac agaaccagtg agtggtgaga gctctgtcag 180
tgacaaacac tcctttggcc tgttgaattt gctgaagaac atcacctaaa gtctgcacac 240
gagcccatth ttaccaagat ttgatcagtg tctttactga gctggaagcc tctgaaagtt 300
attaaaggac agaatccaaa agaatgcctt taattcttgt ctgagaatct tgg 353

```

<210> 1179

<211> 288

<212> DNA

<213> Homo sapien

<400> 1179

```

ccaatgggat cctcaagggt cctgccatca atgtcaatga ctccgtcacc aagagcaagt 60
ttgacaacct ctatggctgc cgggagtccc tcatagatgg catcaagcgg gccacagatg 120

```



```
<210> 1180
<211> 523
<212> DNA
<213> Homo sapien
```

```
<210> 1181
<211> 493
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(493)
<223> n = A,T,C or G
```

```
<210> 1182
<211> 329
<212> DNA
<213> Homo sapien
```

[illegible]

<210> 1183
 <211> 198
 <212> DNA
 <213> Homo sapien

<400> 1183
 cctgacagac agaagggcctt ggagatTTTT tttctttaca attcagtctt cagcaacttg 60
 agagctttct tcatgttgct aagcaacaga gctgtatctg caggttcgta agcatagaga 120
 cgatttgaat atcttccagt gatatcggt ctaactgtca gagatgggtc aacaaacata 180
 atcctgggga catactgg 198

<210> 1184
 <211> 224
 <212> DNA
 <213> Homo sapien

<400> 1184
 ctggaggtgc ctcagaaggt gcattctgct tcttgcaggg gcttgaaaca ccaaggcact 60
 ccagggatcc tggagtcaaa gcagcagccc cgggtgttgct actccttggg ggtgacatgg 120
 gggtagccgc agtccaccct gtccttggct ggcacggcac actggtttgc agacaggccc 180
 acgtactcct cagcagagct ggaggacagc aaggccagga ccag 224

<210> 1185
 <211> 367
 <212> DNA
 <213> Homo sapien

<400> 1185
 cctttttacag atgtcagctt tctactggcct ccatgcacaa cctcccacta ccacccaatc 60
 tgcctgcccac agcaaagtgc aggcaccctg ggccccctgg aggatgcggg caggggctac 120
 agggcatcca ggatgtggct gatcttgggt accagctcct ggcgctttcc tgagatgagc 180
 ttctcattct caatgtacgt gtctttcttg agcttgccag ccaccaggcg ctcagcctcc 240
 accgccgact tcagcaccag ctcttggacc tgtgcatcca gcttctgcat ttcgtcact 300
 ctgtcgcaca gatcagagcc ctctgtcttc agcctggact gcagcagtgc aatctcactg 360
 gtcaagg 367

<210> 1186
 <211> 188
 <212> DNA
 <213> Homo sapien

<400> 1186
 ccattaagcg gatgctggag atgggagcta tcaagaacct cacgtccttc cgacctgggc 60
 aagagctgta gcctgtcggg tgcctactct gctgtctggg tgacccccat gcgtggctgt 120
 gggggtggct ggtgccagta tgaccactt ggactcacc cctcttgggg agggagtcc 180
 gggcctgg 188

<210> 1187
 <211> 379
 <212> DNA
 <213> Homo sapien

<400> 1187
 gttgatgcta ctctgaagtc tctcaacaac cagattgaga cccttcttac tcttgaaggc 60

```

tctagaaaaga gccagctcg cacatgccgt gacttgagac tcagccaccc agagtggagc 120
agtggttact actggattga ccctaaccaa ggatgcacta tggatgctat caaagtatac 180
tgtgatttct ctactggcga aacctgtatc cgggcccacac ctgaaaacat cccagccaag 240
aactgggtata ggagctccaa ggacaagaaa cacgtctggc taggagaaac tatcaatgct 300
ggcagccagt ttgaatataa tgtagaagga gtgacttcca aggaaatggc tacccaactt 360
gccttcatgc gcctgctgg 379

```

<210> 1188

<211> 384

<212> DNA

<213> Homo sapien

<400> 1188

```

cgcgtcggac tgcagccagt ccgtttcctt tctttagcca gccatcctgg tactgtagtt 60
taggggttga tgggtggtga aattgatttc tggctgggta ctaagggtgcc tgctagccat 120
tgtataaaat taaaacatga agaataattt ttttttgagc atggctagtg gatttaaaac 180
aacacatacc tgtcactgct ggagtcaaac ttataaaaag ccttaagtgg aaagtgttcc 240
agacggagac tctgagttaa tagaggagta gaagctggtg tttaaagttcc cacgacgcac 300
atggctttgc cagaaactct gtttaatgat cggcctttca cctcttcact tatecttagt 360
cccagtagcc aggatacctg atgg 384

```

<210> 1189

<211> 419

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(419)

<223> n = A,T,C or G

<400> 1189

```

ggaaaaacca gccactgctt tacaggacag ggggttgaag ctgagccccg cctcacaccc 60
acccccatgc actcaaagat tggattttac agctacttgc aattcaaaat tcagaagaat 120
aaaaaatggg aacatacaga actctaaaag atagacatca gaaattgttg agttaagctt 180
tttcaaaaaa tcagcaattc cccagcgtag tcaagggtgg aactgcacg ctctggcatg 240
atgggatggc gaccgggcaa gctttcttcc tcgagatgct ctgctgcttg agagctattg 300
ctttgttaag atataaaaag gggtttctt ttgtctttct gtaaggtnna cttccagctt 360
ttgattgaaa gtcttagggg gattctatct ctgctgtgat ttatctgctg aaagctcag 419

```

<210> 1190

<211> 173

<212> DNA

<213> Homo sapien

<400> 1190

```

ccaggtactg gcacatcatg ctctggatgg ggggtggtgg gtctgttagg cagagaaaca 60
ggaaattgtc gtagtcagta tcgagcagc tggcctcggt cggcaccgta tagttgatct 120
tgaacttctt tggattctca gtcttctctc caaggacctt cttctcaaca cag 173

```

<210> 1191

<211> 341

<212> DNA

<213> Homo sapien

<400> 1191
 cctcctgcca gcagttcttg aagcttcttt ttcattcctg ctactctacc tgtattttctc 60
 agttgcagca ctgagtgggc aaaatacatt tctggggccac ctcaggggaac ccatgcatct 120
 gcctggcatt taggcagcag agcccttgac cgtccccccac agggctctgc ctcacgtcct 180
 catctcattt ggctgtgtaa agaaatggga aaagggaaaaa ggagagagca attgaggcag 240
 ttgaccatat tcagttttat ttattttatt ttaatttggt cttttctcca agtccaccag 300
 tctctgaaat tagaacagta ggcggtatga gataatcagg a 341

<210> 1192

<211> 324

<212> DNA

<213> Homo sapien

<400> 1192
 ttggaggttg gcggcgcggg gctgaaggct agcaaaccga gcgatcatgt cgcacaaaaca 60
 aatttactat tcggacaaat acgacgacga ggagtttgag tatcgacatg tcatgctgcc 120
 caaggacata gccaaagtgg tccctaaaac ccatctgatg tctgaatctg aatggaggaa 180
 tcttggcggt cagcagagtc agggatgggt ccattatatg atccatgaac cagaacctca 240
 catcttgctg ttccggcgcc cactacccaa gaaaccaaag aaatgaagct ggcaagctac 300
 ttttcagcct caagctttac acag 324

<210> 1193

<211> 521

<212> DNA

<213> Homo sapien

<400> 1193
 ctgctttgtt ttctgttggc agtggaggga caagggtgaga ggagccaggg gtagtcatga 60
 acaccagtgg gttctgcctt gggcagctcc ccaccttctt taagagagta ctgtgtctca 120
 gctccagcag tctcaactgg gaagaccag gactcctgct cttttctcta atccctggga 180
 gacgaggtcc agctaaggta gagtaagcag tcagtgaacca ggcaggctgg tttgggaggt 240
 cactgcctgg aggacgggat cttgtattct tcggaagatg gctgggaaat tcttccctcc 300
 attacgtaga actttcttcc cctcctcagt tgagggtgcct agatgtccca caacgggggtc 360
 ttcactcagg tcctccagag gcacacgctc aaacagtggg tgctcttcga aatgagtga 420
 catccagtcg tgtagctcca gcacatcggg tatggtatac accagcccct gcataggcaa 480
 aatcaccccta gacaggaggc tgcattgcaac gtcagcagcc a 521

<210> 1194

<211> 208

<212> DNA

<213> Homo sapien

<400> 1194
 ccagtgacta gaaggcgagg cgccgcggga ccatggcggc ggcggcggac gaggcgagtc 60
 cagaggacgg agaagacgag ggagaggagg agcagttggg tctggtggaa ttatcaggaa 120
 ttattgattc agacttcctc tcaaaatgtg aaaataaatg caagggtttg ggcattgaca 180
 ctgagaggcc cattctgcaa gtggacag 208

<210> 1195

<211> 499

<212> DNA

<213> Homo sapien

<400> 1195
 ccagaaagga aagacaataa ttttgttttt tcattttgaa aaaattaaat gctctctcct 60
 aaagattctt cacctacttt ggtctccata acttctatgt tttctttcct tctgacacac 120
 tagtgcccct aaattgtgat ttgcctatac gtttagggcc ggggttgga gatgttaaca 180
 accatttaag attcatttct gcagtgggag tgggtggagt ttcacctctt gggaaagggg 240
 caggtgacag gtattttatca gtcagtgcct ctctagctct tgtaggaaga agcacacgca 300
 ggatggagtc tagaggatga gcgatattga ctagcaattc atgggctccc tccagcagtg 360
 cgagggtcag agtttctgga gccttgggag gaggcattcc tgtgaggggg ggtagggag 420
 atgggagggc accaggaaaa gtgattagaa gtcagggtat ggaaggctaa attaggacag 480
 agtcgagtac atctctgct 499

<210> 1196

<211> 455

<212> DNA

<213> Homo sapien

<400> 1196
 ctgaccccc tttgtccaca gctaagatgg cagcagaatg ctatgtcact atatacagaa 60
 acaagacaac ctgaagctaa atggatgcc cctgcagagt caacagggtcc agcctcacag 120
 tgcacgccct gagctacagc ctctcccaaa aggcattctt cccacagcct caacgccgag 180
 caaggagcat caagggtttg tctcggttgt tttgttcttt ttacaaacta tagatatata 240
 cagttgaaaa ctcaggattt ctagccaata accatagtta ccaccaacct acaataaaaa 300
 agaaaatgcc agaaacatct ttaaatgcct tgtcacacca acagcaaagt gcacagagtg 360
 aggagaacac gagagtgcct tttcatttta aaaatgtttg gaaatatgta caacttcgat 420
 acagtttcag ggtgctccag acacccatgg acctg 455

<210> 1197

<211> 444

<212> DNA

<213> Homo sapien

<400> 1197
 cctggatgtg gctcttcgca ctgaaggcca agtagtagat cacaaggccg atcgccgcag 60
 ccagcacctc agtggacacc cagggcccgt tccaagtgcc ccgatgggtc acgtgactg 120
 taaacagagg cgggatgatg gaaatgtcct cgttattcct ctgagccttc ctgaggaggc 180
 tgtaggactc ctcgtcgaag aatctaacct cataggtgcc tgcgtgggag ctcttgtggt 240
 tcaggcttca ggacacctga taacgcccc catcctggcc tcgagtgaca gggaattgtt 300
 ttccaccgac gtcagcatag agagccatgt tctggacctt gttcttgcat gtcagggaga 360
 tctccacaat gaagacggtc tcagtggaaa tgacagcgtc agaagtgggtg tagtaggaag 420
 gggtagctg gggctccagg cagg 444

<210> 1198

<211> 450

<212> DNA

<213> Homo sapien

<400> 1198
 ccatgggtgt ctggagcacc ctgaaactgt atcaaagttg tacatatattt caaacatttt 60
 taaaatgaaa aggcaactct gtgttctcct cactctgtgc actttgctgt tgggtgagca 120
 aggcatttta agatgtttct ggcattttct ttttatttgt aagtggtgtg taactatggt 180
 tattggctag aaatcctgag ttttcaactg tatatatcta tagtttgtaa aaagaacaaa 240
 acaaccgaga caaacctttg atgctccttg ctcgcggttg aggctgtggg gaagatgcct 300
 tttgggagag gctgtagctc agggcgtgca ctgtgaggct ggacctgtt actccgcagg 360
 gggcatccat ttagcttcag gttgtcttgt ttctgtatat agtgacatag cattctgctg 420

ccatcttagc tgtggacaaa ggggggtcag

450

<210> 1199

<211> 294

<212> DNA

<213> Homo sapien

<400> 1199

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| agtcacagtt | gcacctattc | aaaactagct | ttaaagtgag | ctatttttaa | acttcataaa | 60 |
| aatattcatg | attttattag | tttgaatatt | tctacaagat | tcgggtgggc | ttttccttta | 120 |
| ggtgaaaaca | gctatccact | cctgtggcct | tataactcag | gaaatgctgg | ggatgcaaac | 180 |
| gtgcaaaagg | cagggggaag | ctgccaggc | tgagactgga | gcagctagga | gtgtgcttgg | 240 |
| ggaacgggag | ctgagatccc | ggagcagaaa | tggtcagccg | tgctctggag | cagg | 294 |

<210> 1200

<211> 258

<212> DNA

<213> Homo sapien

<400> 1200

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| agctacctaa | gaacagctaa | aagagcacac | ccgtctatgt | agcaaaatag | tgggaagatt | 60 |
| tataggtaga | ggcgacaaac | ctaccgagcc | tggtgatagc | tggttgcca | agatagaatc | 120 |
| ttagttcaac | tttaaatttg | cccacagaac | cctctaaatc | cccttgtaaa | tttaactgtt | 180 |
| agtccaaaga | ggaacagctc | tttggacact | aggaaaaaac | cttgtagaga | gagtaaaaaa | 240 |
| tttaacaccc | atagtagg | | | | | 258 |

<210> 1201

<211> 403

<212> DNA

<213> Homo sapien

<400> 1201

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ctgagctgct | gtctgctttg | gaaaaccggt | cctgccgctg | ccgatggatg | gaaatgcaat | 60 |
| ggatttcagc | ttcttatcat | cagccagggc | caagcagttt | ttcactgtct | ttccagaag | 120 |
| ttcttcacac | ttgtctgcac | cccaaactgg | actattacag | tggtacacaa | acttggcagg | 180 |
| caggccatgg | cctgcgctga | cagcagctcc | agctacttcc | aagggcccgt | tctttttccg | 240 |
| gagttccagg | acagcttcca | caaactcctt | gccacctttc | ttctccagcg | tgtttccctag | 300 |
| gtcatcttta | aggtcaatgt | cagcattggt | aggattgatt | atggcctcca | cctcaaagcc | 360 |
| ggctaaatta | ctgatttcac | tgtgaataag | gttcggcttc | tgg | | 403 |

<210> 1202

<211> 325

<212> DNA

<213> Homo sapien

<400> 1202

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ctgaacctgc | gggagtcggc | caccatcacg | tgcttggtga | cgggcttctc | tcccgcggac | 60 |
| gtcttcgtgc | agtggatgca | gagggggcag | cccttgctcc | cggagaagta | tgtgaccagc | 120 |
| gccccaatgc | ctgagcccca | ggccccaggc | cggtaacttcg | cccacagcat | cctgaccgtg | 180 |
| tccgaagagg | aatggaacac | gggggagacc | tacacctgcg | tggtggccct | tgaggccctg | 240 |
| cccaacaggg | tcaccgagag | gaccgtggac | aagtcaccgc | gtaaaccac | cctgtacaac | 300 |
| gtgtccctgg | tcatgtccga | cacag | | | | 325 |

<210> 1203

<211> 518
 <212> DNA
 <213> Homo sapien

<400> 1203
 ctcaaccaca gtctgacacc agagcccact tccatcctct ctggtgtgag gcacagcgag 60
 ggcagcatct ggaggagctc tgcagcctcc acacctacca cgacctccca gggctgggct 120
 caggaaaaac cagccactgc tttacaggac aggggggttga agctgagccc cgcctcacac 180
 ccacccccat gcactcaaag attggatttt acagctactt gcaattcaaa attcagaaga 240
 ataaaaaatg ggaacataca gaactctaaa agatagacat cagaaattgt taagttaagc 300
 tttttcaaaa aaccagcaat tccccagcgt agtcaagggg ggacactgca cgctctggca 360
 tgatgggatg gcgaccgggc aagctttctt cctcgagatg ctctgctgct tgagagctat 420
 tgctttgtta agatataaaa aggggtttct ttttgtcttt ctgtaagggtg gacttccagc 480
 ttttgattga aagtcctagg gtgattctat ttctgctg 518

<210> 1204
 <211> 352
 <212> DNA
 <213> Homo sapien

<400> 1204
 ggggaaagga ggtctcactg agcaccgtcc cagcatccgg acaccacagc ggccttctgc 60
 tccacgcaga aaaccacact tctcaaacct tcaactcaaca ctcccttccc caaagccaga 120
 agatgcacaa ggaggaacat gaggtggctg tgctgggggc acccccagc accatccttc 180
 caaggtccac cgtgatcaac atccacagcg agacctccgt gcccgaacct gtcgtctggt 240
 ccctgttcaa caccctcttc ttgaactggt gctgtctggg cttcatagca ttgcctact 300
 ccgtgaagtc tagggacagg aagatggttg gcgacgtgac cggggcccag ga 352

<210> 1205
 <211> 250
 <212> DNA
 <213> Homo sapien

<400> 1205
 ctgttcaact tccaactcta aataggcacc attaaacaaa aaacccagc attttaaatt 60
 tctccagcac acattccagg atcaatgctc tgaactgtaa tcagctagta attcataacg 120
 ggaatacagc cttagaatgg aagctatatt gcttccctgc cccctttctc ttacaattgg 180
 agagtgtagg tattaaggga taaaaagtca gaggaagaat aattaaaaag aaaaatgccc 240
 aaagctgcag 250

<210> 1206
 <211> 275
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(275)
 <223> n = A,T,C or G

<400> 1206
 ctgctctcgn ngnetcactg gatggaccag cacttccgca cgacgcccct ggagaagaac 60
 gccccgctct tgctggccct gctgggtatc tggatcatca actgcttttg gtgtgagaca 120
 caogccatgc tgccctatga ccagtacctg caccgctttg ctgcgtactt ccagcagggc 180

gacatggagt ccaatgggaa atacatcacc aaatctggaa cccgtgtgga ccaccnnaca 240
ggccccattg tgtgggggga gccagggacc aatgg 275

<210> 1207
<211> 182
<212> DNA
<213> Homo sapien

<400> 1207
ccatctcctg ctgcaagtcc agggcgacgt agcacagctt ctctttgatg tcgcgcacga 60
tttcccgctc ggccgtggtg gtgaagctgt agcctcgctc agtgaggatc ttcattgaggt 120
agtcggtcag gtcccgcca gccaggtcca gacgcaggat ggcgtggggg agggcgtagc 180
cc 182

<210> 1208
<211> 260
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(260)
<223> n = A,T,C or G

<400> 1208
gctggttatg aactcctgac ctcaagtgat ctgccctcct cagcctccca aagtgtctggg 60
attataggca tgagccactg gaatttttct tttttttttt ctttcttttt tttttttttt 120
ttaaattgan acaaggtctg gctctatcgc ccangctgga gtgcagnggc accatntcgg 180
ctcactgcaa cctctgcctg ctgggctcga gccatcctcc cacctcagcc tcccaagtan 240
ttgggactag aggtatgcac 260

<210> 1209
<211> 487
<212> DNA
<213> Homo sapien

<400> 1209
aaaccactc caccttacta ccagacaacc ttagccaaac catttaccca aataaagtat 60
aggcgataga aattgaaacc tggcgcaata gatatagtag cgcaagggaa agatgaaaaa 120
ctataaccaa gcataatata gcaaggacta atccctatac cttctgcata atgaattaac 180
tagaaataac tttgcaagga gagccaaagc taagaccccc gaaaccagac gagctaccta 240
agaacagcta aaagagcaca cccgtctatg tagcaaaata gtgggaagat ttataggtag 300
aggcgacaaa cctaccgagc ctggtgatag ctggttgctc aagatagaat cttagttaa 360
ctttaaattt gccacagaa ccctctaaat ccccttgtaa atttaactgt tagtccaaag 420
aggaacagct ctttggacac taggaaaaaa ccttgtagag agagtaaaaa atttaacacc 480
catagta 487

<210> 1210
<211> 216
<212> DNA
<213> Homo sapien

<400> 1210
ccactcagct cagcggggcga cgtgccccta caagttggca gaagtggctg ccactgctgg 60


```

gtttgtgtaa gagagggtgc tgccaccatt acctgcagaa accttctcat aggggctacg 120
atcgggtactg ctaggggggca catagcgccc atggatgtgg taggtggggg actcgtcat 180
aggatggtag gtatcccggg ctggaaagat gtccag 216

```

<210> 1211

<211> 443

<212> DNA

<213> Homo sapien

<400> 1211

```

ccaaggtcag aggtgatgc aacaggccct cttctcccca gggccaggct cctgtccagc 60
ctgggcactg cccagagtga tggcattggg cgggatgctg ttctgtctct gcttggacac 120
cttcgcaaag atttctttca ggacagtctc aaaggctagc tcaacattgg tagagtccag 180
ggctgaggtc tccaggaaga gcagtcatt gttttcagcg aacattcggg cctcctcagt 240
gggcacttcc cgggcctggc tgaggctcact tttgttacct acgagcatga cgacgatcgt 300
ggcttcagca tggcataga gctccttcag ccacgtctcc accacagcat aggtctgggtg 360
cttggttagg tcaaacacca ggagggcccc cactgcacca cgatagtacg ccgaggtgat 420
ggctcggtac cgctccaggc cag 443

```

<210> 1212

<211> 526

<212> DNA

<213> Homo sapien

<400> 1212

```

actgaaaccc gagtaaagtc tcaggctgct gcatatgaat acatggctgc atacatagaa 60
aatgcgaaac aggttggccg ccttgaaaat gcaatcgggt ggtatcatag ccaccctggc 120
tatggctgct ggctttctgg gattgatgtt agtactcaga tgetcaatca gcagttccag 180
gaaccatttg tagcagtggg gattgatcca acaagaacaa tatccgcagg gaaagtgaat 240
cttggcgctt ttaggacata cccaaagggc taaaaacctc ctgatgaagg accttctgag 300
taccagacta ttccacttaa taaaatagaa gattttgggtg tacactgcaa acaatattat 360
gccttagaag tctcatattt caaatcctct ttggatcgca aattgcttga gctgttgggtg 420
aataaatact ggggtgaatac gttgagttct tctagcttgc ttactaatgc agactatacc 480
actggtcagg tctttgattt gtctgaaaag ttagagcagt cagaag 526

```

<210> 1213

<211> 359

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(359)

<223> n = A,T,C or G

<400> 1213

```

ccagccattg cctgncattt ggtagtatag tatgattctc accattattt gtcattggagg 60
cagacataca ccagaaatgg gggagaaaca gtacatatct ttctgtcttt agtttattgt 120
gtgctggtct aagcaagctg agatcatttg caatggaaaa cacgtaactt gtttaaaagt 180
ttttctggta gcttttagctt tatgctaaaa aaaataatga cattgggtat ctatttcttt 240
ctaagactac attantanga aaataagtct tttcatgctt atgatttagc tgttttgtgg 300
taattgcttt ttaaaggaag nnattaatat cataagttat tattaatatt gtgaacnca 359

```

<210> 1214

<211> 428
 <212> DNA
 <213> Homo sapien

<400> 1214
 ccaagcttga ggcagcccta ggtgaggcca agaagcaact tcaggatgag atgctgcggc 60
 ggggtggatgc tgagaacagg ctgcagacca tgaaggagga actggacttc cagaagaaca 120
 tctacagtga ggagctgcgt gagaccaagc gccgtcatga gacccgactg gtggagattg 180
 acaatgggaa gcagcgtgag tttgagagcc ggctggcggg tgcgctgcag gaactgcggg 240
 cccagcatga ggaccagggtg gagcagtata agaaggagct ggagaagact tattctgcca 300
 agctggacaa tgccaggcag tctgctgaga ggaacagcaa cctgggtggg gctgcccacg 360
 aggagctgca gcagtcgcgc atccgcacgc acagcctctc tgcccagctc agccagctcc 420
 agaagcag 428

<210> 1215
 <211> 414
 <212> DNA
 <213> Homo sapien

<400> 1215
 ctgaagcact cttcagagac tacgtccaca gacactgatg ctgaggcctt tcttgtaagt 60
 gaagaaaaag gaatgcagca aagaagagtt cgacattgga gtccttagtt ccatcaggat 120
 cccattcgca gccttttagca tcatgtagaa gcaaactgca cctatggctg agatagggtgc 180
 aatgacctac aagattttgt gttttctagc tgtccaggaa aagccatctt cagtcttgct 240
 gacagtcaaa gagcaagtga aaccattttcc agcctaaact acataaaagc agccgaacca 300
 atgattaaag acctctaagg ctccataatc atcattaaat atgcccacac tcattgtgac 360
 tttttatttt atatacagga ttaaaatcaa cattaaatca tcttattttac atgg 414

<210> 1216
 <211> 162
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(162)
 <223> n = A,T,C or G

<400> 1216
 cctggccgca ggggtccccg gtattgctgt tgctacgagg ttgggggggca gcgattgtcc 60
 tgtgggagcc accgttctcc tgggtcgggg accctcactt cttctggggg gtgctcannt 120
 tctgcatgcc ccgatcttg tccagcangc cagaaatgaa gg 162

<210> 1217
 <211> 392
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(392)
 <223> n = A,T,C or G

<400> 1217

```

ctgaagtaga ggctggaact gaagctgaga ctgaggctga ggctgaaact ggagctaagg      60
gtgaggctgg aactggagct gaggttgagg ccagaactgg agctaaagtt gaggctggaa      120
ccggagctga ggttgaggct ggaactggag ttaaggttgc tggaaagtga gctgaggttg      180
aggctggaac tgaagctgag gttgaagggt gaagtggagc cgaagctaga ggtggaactg      240
aggctgaaga ctgtgcttgc tggatccctg tagcctgttt tttggcaaact cttggaggaa      300
gcttanaagt ctggcttctt cctttttcat ttgcattctt tttgttccag accttaaaaa      360
attaacgggg accatttttg tcaataatgc ag                                     392

```

<210> 1218

<211> 526

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(526)

<223> n = A,T,C or G

<400> 1218

```

ctgagctttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa      60
agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatata ttaacaaagc      120
aatagctctc aagcagcaga gcatctcgag gaagaaagct tgcccggtcg ccatcccatc      180
atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa      240
agcttaactt aacaatttct gatgtctatc ctttagagtt ctgtatgttc ccatttttta      300
ttcttctgaa ttttgaattg caagtagctg taaaatccaa tctttgagtg catgggggtg      360
gggtgtgaggc ggggctcanc ttcaaccccc tgtcctgtaa agcagtggct ggtttttctt      420
gagcccagcc ctgggaggtc gtggtangtg tggaggctgc agagctcctn cagatgctgc      480
cctcgctgtg cctcacacca nagaggatgg aagtgggctc tgggtgt                                     526

```

<210> 1219

<211> 382

<212> DNA

<213> Homo sapien

<400> 1219

```

ctggccggcg gtgcagatct ggagtccagc ctcagggatg cgctactttc cattctctgc      60
attgaacatt cgttctgtca gcatccgctc cagcttccact gcatcagcgg caaacttgcg      120
gatcccgctc gagagcttct ccacagccat ctggtcctcg ttgtgcaacc aacggaaaaga      180
cttctcatcc aggtggattt tttccaggtc actggccttg gccgccttgg ctgagagcac      240
aggcaccagc ttggcgttgt cctgcagcag ctctcccagg agcttgggtg agatggtgag      300
gaagtcacag ccggccagtg ctttgatctc gcccggtgtg cggaaggagg cgcccatgac      360
aatggttttg tagctaaact tc                                     382

```

<210> 1220

<211> 127

<212> DNA

<213> Homo sapien

<400> 1220

```

tcgacctcct tgaagcagac caagtatagc aagcctctaa aaggactact gagaaacaga      60
atcagaaaact ctagaactct agttagggcc cttcagcagg gctgcagagc ctccctggat      120
accaggg                                           127

```

<210> 1221

<211> 304
 <212> DNA
 <213> Homo sapien

<400> 1221
 ccaccccgga gatgacacga ggctcacatg actctagaca cttgggtggaa agtgaggcga 60
 gaaaaacaat gacttgggcc aattacacga ctgcaaagct agagctgcca acagggctcc 120
 agggagcttg gcttctgtag aagttctaag gaagcggtag gaactccacg gcggtggggc 180
 gctaactagc agggaccctt gcaagtgttg gtcggggggc tcgggctgcc tgagctgaca 240
 cgaggggagg ggtctgtgta gccaacaggt gaccgaaggg cttgcctgcc cacagcttac 300
 ttgg 304

<210> 1222
 <211> 309
 <212> DNA
 <213> Homo sapien

<400> 1222
 ctgtcgact cgtagctgca actcaactcaa cttgtcttta gcagcaattt ctgcatagtc 60
 attggcatgt tcacctacct ggatgtccgg gtgaactctc agcatgcctc cagcaaagag 120
 ggagaacttg gtggaattgg agtgaagaca gatctggtgc tcaccagggg tatgggaagt 180
 gaaagtgaac ctgccctcgg agccatactg ccggggccagg atgaccttgt cctctgggtc 240
 ctccacctcc acaaacatgc caagccccgg ggtggccggc tgggtactcct cccgctgctt 300
 gtcatacag 309

<210> 1223
 <211> 390
 <212> DNA
 <213> Homo sapien

<400> 1223
 cctggcctgg gagccctgtg cctactagaa gcacattaga ttatccattc actgacagaa 60
 caggtctttt ttgggtcctt cttctccacc acgatatact tgcagtcctc cttcttgaag 120
 attcttttggc agttgtcttt gtcataaccc acaggtgtag aaacaagggt gcaacatgaa 180
 atctctgttt cgtagcaagt gcatgtctca cagttgtcag tctgccactc cgagtttatt 240
 ggtgtttgtt tcctttgaga tccatgcatt tcctggttga atctcctgga actccctcat 300
 taggtatgaa atagcatgat gcattgcata aagtcacgaa ggtggcaaag atcacaacgc 360
 tgcccaggag aacattcatt gtgataagca 390

<210> 1224
 <211> 407
 <212> DNA
 <213> Homo sapien

<400> 1224
 ccttatgact acaacggccc acgagaaaaa tatggaatcg ttgattacat gatcgagcag 60
 tccgggcctc cctccaagga gattctgacc ctgaagcagg tccaggagtt cctgaaggat 120
 ggagacgatg tcatcatcat cggggtcttt aagggggaga gtgaccacgc ctaccagcaa 180
 taccaggatg ccgctaacaa cctgagagaa gattacaaat ttaccacac tttcagcaca 240
 gaaatagcaa agttcttgaa agtctcccag gggcagttgg ttgtaatgca gcctgagaaa 300
 ttccagtcca agtatgagcc ccggagccac atgatggacg tccagggctc caccaggagc 360
 tcggccatca aggacttcgt gctgaagtac gcctgcccc ttggttgg 407

<210> 1225

<211> 250
 <212> DNA
 <213> Homo sapien

```
<400> 1225
ctgcagcttt gggcattttt ctttttaatt attcttcctc tgacttttga tcccttaata      60
cctacactct ccaattgtaa gagaaagggg gcaggggaagc aatatagctt ccattctaag      120
gctgtattcc cgttatgaat tactagctga ttacagttca gagcattgat cctggaatgt      180
gtgctggaga aatttaaaat actgggggtt tttgtttaat ggtgcctgtt tagagttgga      240
agttgaacag                                     250
```

<210> 1226
 <211> 444
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(444)
 <223> n = A,T,C or G

```
<400> 1226
ccttttagct gttgctctgg gcaggggggtg ggggtgcggg ggcttacagt gggggccctt      60
agttggcaca ggttcggaag ggcccagggc agacatgaat tctcctgaga cttgaggtag      120
gttgcttcag ccagcccggg cggagaagaa gggcagagag cgaacatagg agtccagtcg      180
ggagcgaaag agctcacttt gcacagtttg gcccagcggg cacaggggat tcttcaccac      240
cagctccaca tacagcgcac tgtagatgtg gtgcagcaca tctcggatgg gtcccacgcc      300
caagtcagta ttcataaaca ctttgatccc agtgggcgtc tcgtagtaat ggagtttgta      360
acggctagtt tggaaggcca ggaagccatc cttcatgtct agcggggaca tcttgctgac      420
aaacgancgg atagagaaga gcat                                     444
```

<210> 1227
 <211> 491
 <212> DNA
 <213> Homo sapien

```
<400> 1227
gttagcctta catgttgtgt agacttactt taagtttgca cccttgaaat gtgtcatatc      60
aatttctgga ttcataatag caagattagc aaaggataaa tgccgaaggt cacttcattc      120
tggacacagt tggatcaata ctgattaagt agaaaatcca agctttgctt gagaactttt      180
gtaacgtgga gagtaaaaag tatcggtttt attctttgct gatgtccttt ctgcttgaaa      240
taacagtcac catacagcta aaggagagga gtttctttcc ttctaagtag gcagaaatgg      300
tatcattatg ttgccgctct ccaatctccc agagctcgtt ctctagagaa tcaccttctt      360
tcgctttttt tttttttttg aggtagagtc tcactatgtt gcccagacta gccttgaaact      420
cctgggctca agtgattctc cctcctcagc ctcccagagta gctggaacga actatagttg      480
caccactgca g                                     491
```

<210> 1228
 <211> 279
 <212> DNA
 <213> Homo sapien

```
<400> 1228
ctgggcggat ctgatcaact aggcaacatc atgtccggat atgagttcat caacaagttg      60
```


$\langle 400 \rangle$ 1236

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgatcctca | ctattgtggg | caccatcgct | ggcatcgta | ttctcagcat | gataattgca | 60 |
| ttgattgtca | cagcaagatc | aaataacaaa | acgaagcata | ttgaagaaga | gaacttgatt | 120 |
| gacgaagact | ttcaaaatct | aaaactgcgg | tcga | | | 154 |

<210> 1237

<211> 375

<212> DNA

<213> Homo sapien

<400> 1237

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ccactggatc | tttgggatta | aagctctgtt | ggatttgtac | ctcagaggaa | gatcaagtgg | 60 |
| ctgatccttt | ggactctgta | aagagcattc | ttctagtcag | aggggtggaat | ggcagcagca | 120 |
| actggaagaa | aatgagtttt | ttggtgcccc | cacccaagag | cacacacatg | ctgcactgtc | 180 |
| tcggaagca | gggccagcta | gagccaccat | gttcttcctt | acctcagttt | acctgcggcc | 240 |
| tgcgctgcac | tgcagatgcc | caccctgccc | tgggtctggc | cggcggaagc | tctgtccaag | 300 |
| gtccacacac | ctccaggttt | acgccaacat | ccttgtgccc | tccccacctt | ctcttccaac | 360 |
| gcattaggtg | cattg | | | | | 375 |

<210> 1238

<211> 454

<212> DNA

<213> Homo sapien

<400> 1238

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| gtcaagatca | agttcaatat | catcgctctt | ctctatgact | acaaccccaa | cctggcaacc | 60 |
| tacatgaagc | cagagatgtg | ggggaagtgc | ctggactgca | tcaatgagct | gatggatata | 120 |
| ctgtttgcaa | atcccaacat | ttttgttggg | gagaatattc | cgggaagagag | tgagaacctg | 180 |
| cacaacgctg | accagccact | gcgtgtccgt | ggctgcattc | taactctggt | ggaacgaatg | 240 |
| gatgaagaat | ttaccaaaat | aatgcaaaat | actgaccctc | actccaagag | tacgtggagc | 300 |
| acttgaagga | tgaggcccag | gtgtgtgcca | tcatcgagcg | tgtgcagcgc | tacctggagg | 360 |
| agaagggcac | taccgaggag | gtctgccgca | tctacctgct | gcgcattcctg | cacacctact | 420 |
| acaagtttga | ttacaaggcc | catcagcgac | agac | | | 454 |

<210> 1239

<211> 483

<212> DNA

<213> Homo sapien

<400> 1239

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ctgccaggct | gaaaagaagc | ctcagctccc | acaccgcctt | cctcaccgcc | cttcctcggg | 60 |
| agtcacttcc | actggtggac | cacgggcccc | cagccctgtg | tcggccttgt | ctgtctcagc | 120 |
| tcaaccacag | tctgacacca | gagcccatct | ccatcctctc | tgggtgtgagg | cacagcgagg | 180 |
| gcagcatctg | gaggagctct | gcagcctcca | cacctaccac | gacctcccag | ggctgggctc | 240 |
| aggaaaaacc | agccactgct | ttacaggaca | gggggttgaa | gctgagcccc | gcctcacacc | 300 |
| cacccccatg | cactcaaaga | ttggatttta | cagctacttg | caattcaaaa | ttcagaagaa | 360 |
| taaaaaatgg | gaacatacag | aactctaaaa | gatagacatc | agaaattgtt | aagttaagct | 420 |
| ttttcaaaaa | atcagcaatt | ccccagcgta | gtcaagggtg | gacactgcac | gctctggcat | 480 |
| gat | | | | | | 483 |

<210> 1240

<211> 358

<212> DNA

<213> Homo sapien


```
<210> 1241
<211> 194
<212> DNA
<213> Homo sapien
```

```
<210> 1242
<211> 316
<212> DNA
<213> Homo sapien
```

```
<210> 1243
<211> 275
<212> DNA
<213> Homo sapien
```

```
<210> 1244
<211> 235
<212> DNA
<213> Homo sapien
```

| | | | | | | | | | |
|------------|------------|------------|------------|------------|------------|--|--|--|-----|
| <400> | 1244 | | | | | | | | |
| ctgctgcgct | tggataacaa | gtaattcaac | gcacgcactt | aacagaaatg | ttaaactata | | | | 60 |
| acaagcacca | tttgaggatt | aacaggaaca | tttttttgaa | gatttcaaac | gaactcgact | | | | 120 |
| ttcagtataa | tgtacctaa | agtatttata | aacagctcat | cggagcctct | atttgtcata | | | | 180 |
| qacttttgag | ttgattgttg | ggaccacata | ataggacct | tttttttttg | tcttt | | | | 235 |

```
<220>
<221> misc_feature
<222> (1)...(640)
<223> n = A,T,C or G
```

```
<210> 1246
<211> 509
<212> DNA
<213> Homo sapien
```

```
<210> 1247
<211> 310
<212> DNA
<213> Homo sapien
```

<210> 1248

<211> 640
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(640)
 <223> n = A,T,C or G

<400> 1248
 aaagatataa aactatggag aaaactgcta aagggtatcc ctgaccttta tgatgatgca 60
 gctatttttcg aggccaaaaa atcatttttac tgggcaagaa aaacatctca ttcctttgtc 120
 gtgaatatcc ttgctcaggc tctttatgaa ttattttctg ccacagatga ttccctgcat 180
 caactaagaa aagcctgttt tctttatttc aaacttggtg gcgaatgtgt tgcgggtcct 240
 gttgggctgc tttctgtatt gtctcctaac cctctagttt taattggaca cttctttgct 300
 gttgcaatct atgccgtgta tttttgcttt aagtcagaac cttggattac aaaacctcga 360
 gcccttctca gtagtggtgc tgtattgtac aaagcgtggt ctgtaaatatt tcctctaatt 420
 tactcagaaa tgaagtatat gggtcattaa gcttaaaggg gaaccatttg tgaatgaata 480
 tttggaactt accaagtcct aagagacttt tggaagagga tatatatagc atagtaccat 540
 accacttata aagtggaaac tcttggaacca agatttggtat taatttggtt ttgaagtttt 600
 tggnatataa atatgtaaat acatgcttta attgcaattt 640

<210> 1249
 <211> 1108
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(1108)
 <223> n = A,T,C or G

<400> 1249
 caaaataaat ttcaattcaa tgaaaagtaa ataacttagg gatctataaa tgacactgca 60
 atgtatcttg ttccattttt aacaggaagt ctttcatgca aatgtgtgag tctcccagga 120
 tgcataagc tccagccttt tcgtggtgac tcaatagagc aattgtacct taaaaatktg 180
 caaccacctc cctgaaagtc ttctcccacg ttattaagtg caatgyttat ggtaaatgta 240
 gaagcatcat gatgaggacg aagagaacgc tgtcgttcag gggagtattt tactacaaaa 300
 ttcagtagtg caaatccctt cgtataatag cctgcaaaga ctttcagtgt aactgggtgca 360
 atgaactccc ggataaaatg aagccataca ttctccagat caacttgctt catgtggata 420
 tcatcagttg ggacattttt ataaccacca gatatacggc tatcatgatg tttttcccca 480
 gaccatttgc cgtaatgttc catttcttct accaattcat cacaggncct tttcagaaaa 540
 tatggggaac cmaaaagaca tctggacagg gctgttcaam ctatattttc agtgaaaatc 600
 tttgaataat ccmcggttta tatacttttc cttccagtcc acaggatttt caaaaatctg 660
 ccagaggtca ttgttataat gggaagtatt gtaattagca gtggataata gccttccaaa 720
 ttcattgtct ttagaaatgt acataaatac accctttggg gggctgagca tttggaatgt 780
 ttccggagta ggggagtctt tttccctttg taaagtcatt tctctagcat ttccggcaag 840
 agccatatca ggatccagtt tatcacgaac aaaatagctc ctttcattca tctctgatcg 900
 gagtgtcttt cctttaatta agtacacatt agccatatat gggacattcc atactcctac 960
 tctattccct tgaacaatat ccacataatc ttcagatcgt gcatagtatc catcaggact 1020
 caatgtctcc cagaaattgg accacagctt tccatgacga gttacaagag gagcaatgat 1080
 ctttctgttt tgttcaatca aaattttt 1108

<210> 1250

<211> 567
 <212> DNA
 <213> Homo sapien

<400> 1250
 ctgaatattg aactggaagc agcacatcat taggcctttat gactgggtgt gtgttgtgtg 60
 tatgtaatac ataatgttta ttgtacagat gtgtgggggt tgtgttttat gatacattac 120
 agccaaatta tttgttggtt tatggacata ctgccctttc attttttttc tttccagtg 180
 tttaggtgat ctcaaattag gaaatgcatt taaccatgta aaagatgagt gctaaagtaa 240
 gctttttagg gccctttgcc aataggtagt cattcaatct ggtattgatc ttttcacaaa 300
 taacagaact gagaaacttt tatatataac tgatgatcac ataaaacaga tttgcataaa 360
 attaccatga ttgctttatg tttatatatta acttgtatatt ttgtacaaac aagatttgtg 420
 aagatatatt tgaagtttca gtgatttaac agtcctttcca acttttcatg atttttatga 480
 gcacagactt tcaagaaaat acttgaaaat aaattacatt gccttttgtc cattaatcag 540
 caaataaaac atggccttaa ctaaaaa 567

<210> 1251
 <211> 655
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(655)
 <223> n = A,T,C or G

<400> 1251
 gaaagaaacc aattttaatgc caccaaaccat aagcctgcta tacctgggaa acaaaaaatc 60
 tcacacctaa attctagcag agtaaacgat tccaactaga atgtactgta tatccatag 120
 gcacatttat gactttgtaa tatgtaattc ataatacagg ntttaagggtgt gtggnatgga 180
 gctaggaaaa ccnaaggagn aggaaattat nnaaaagaac tgnaggtnaa gtataaagtc 240
 atatgcctga tttcctcaaa ccttttggtt ttccctcatgg cttctggcctt tatattttta 300
 tcacaaacca agatctaaca gggntccttc tagaggatta ttagataagt aacacttgat 360
 cattaagcac ggatcatgcc actcattcat gggtgntcta tgttccatga actctaatag 420
 cccaacttat acatggcact ccaaggggat gcttcagcca gaaagtaaag ggctgaaaaa 480
 gtagaacaat acaaaagccc tcgtgtgggg ggaactgnng gctcactctt acttggcctt 540
 cattcnaaac aggttgggnc tttcntgcga ngatctctca gggnggtaaa aactttntgg 600
 ntttcaacan aanaggtttg gntgaatgat tactcggcng acacctaagg gatcc 655

<210> 1252
 <211> 672
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(672)
 <223> n = A,T,C or G

<400> 1252
 aaantgcaaa aaccagaag accaataatt ctgaaacttg gcatgagtgt gccagtcag 60
 cagcttgcaa agagaggatg tgtcagttac tacaattgct gtactccttt agctgagtcc 120
 ttcaactttc tccttcttgc cagtaaatac tacgttgtaa ttcatatgac tgagatctta 180
 gtatcacagg attttttagct cccatgcctc cttcaaaatt gtttacatgg atttgtttct 240

```

attctctgtgta ggccatatctc caaacacatt cacttctaaa tccaacacaa gtgaaggacc 300
agccaggatg aaacacttca gcaatcattt tgttaaaaat aacatcctgg tcatcaagct 360
aagcataagc acctcttgta taacaattca tcttaaaagc ttaaagtaca ataataaaaa 420
taactgcttg aaaactggaa atgaaataca acagaaaaac tgaagcatta gtaatttttg 480
caagtaacc aggtacagta catttgattt catagagggt gttttctgat gtttaaggag 540
agggtagaag gggtaggaaa acttggcaag gaagatggaa acagcacaac cagttatttt 600
gcttttaata aagtaaattg aatgacagga gtagggaggt gacaaacaca tcnatatata 660
tttttcttat gg 672

```

<210> 1253

<211> 644

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(644)

<223> n = A,T,C or G

```

<400> 1253
ccaaatattt gttagaaact tctggtaact tagatggtct ggaatacaag ttacatgatt 60
ttggctacag aggagtctct tcccaagaga ctgctggcat aggagcatct gctcacttgg 120
ttaacttcaa aggaacagat acagtagcag gacttgctct aattaaaaaa tattatggaa 180
cgaaagatcc tgttccaggc tattctgttc cagcagcaga acacagtacc ataacagctt 240
gggggaaaga ccatgaaaaa gatgcttttg aacatattgt aacacagttt tcatcagtgc 300
ctgtatctgt ggtcagcgat agctatgaca ttataatgc gtgtgagaaa tatgggggtga 360
agatctaaga catttaatag tatcgagaag tacacagaca ccactaataa tcagacctga 420
ttctggaaac cctcttgaca ctgtgttaaa ggttttggag attttaggta agaagtttcc 480
tgttactgag aactcaaagg gttacaagtt gctgccacc ttatcttaga gttattcaag 540
gggatggagt agatattaat accttcaaaa gagattgnag anggcatgaa acaaaaaaatg 600
yggactattg aaaatattgc ctctggtctg gcggagggtt gctc 644

```

<210> 1254

<211> 438

<212> DNA

<213> Homo sapien

```

<400> 1254
aaagggcatt tgaggggagg attattgcta tgaatgaaaa aaatatttta gcttagacta 60
agctacctgc cttcaaaata gtttagggac caccaccata ttttattttg tttttatttt 120
tgaacatttt tctaattgatt tggagagaaa actattttaca aaaattccac atatcagtga 180
tacaattttt tgctgtcacc aattttttat aatagcagag tggcctgttc taagaaggcc 240
atatttttta agttatcttt cagggttaaca tggaaatact ataaagttgg atgtcaaact 300
ttaatatggt ttcagtgttc tctaattttt tggaaatttt gtagacttta cacctggaaa 360
aaaagatttg taaaatcacc ggaacaattg tgtgctttat tttataggta gtgggttatta 420
gtattacatc cccatttt 438

```

<210> 1255

<211> 519

<212> DNA

<213> Homo sapien

```

<400> 1255
caagcacagg ggagtttata gttctgatgt ctttgacatt ttccctggaa cataccaaac 60

```

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| cctagaaaatg | tttccaagaa | cacctggaat | ttggttactc | cactgccatg | tgaccgacca | 120 |
| cattcatgct | ggaatggaaa | ccacttacac | cgttctacaa | aatgaagcat | cttctgagac | 180 |
| tcacaggaga | atatggaatg | tgatctaccc | aatcacagtc | agtgtgatta | ttttattcca | 240 |
| aatatctacc | aaggaatgac | caggagaata | agatcctccg | atgttcgcaa | tggtgtggtg | 300 |
| tcaggaggct | gcctcttaga | caatctccag | atgtactgtg | atgtgagttt | gaaaaagagt | 360 |
| tcctgaagta | ccacatctgg | gagacatgcc | actagctgag | cttcccaaaa | gtctaccaag | 420 |
| agctgaggaa | ttgtatcttc | atccttagca | caaagcacct | taaaaacagt | aaaaggagcc | 480 |
| tctatattcc | agataaatat | agcactgata | aagcgacag | | | 519 |

<210> 1256

<211> 178

<212> DNA

<213> Homo sapien

<400> 1256

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccatgcagga | gttcatgatc | ctcccagtcg | gtgcagcaaa | cttcagggaa | gccatgcgca | 60 |
| ttggagcaga | ggtttaccac | aacctgaaga | atgtcatcaa | ggagaaatat | gggaaagatg | 120 |
| ccaccaatgt | gggggatgaa | ggcggggttg | ctcccaacat | cctggagaat | aaagaagg | 178 |

<210> 1257

<211> 255

<212> DNA

<213> Homo sapien

<400> 1257

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| gggtccactt | gctgccccat | cattgtatca | ccttccttca | atcttttgge | tgccactctc | 60 |
| atgtagggat | ccacggtgag | gaacaaagct | tcaagcagga | cctctccatt | ttttaagggg | 120 |
| gggagctcag | atgtcttcaa | ctcaaagtca | ctattagtag | gatagccaac | aaagtgcctc | 180 |
| ttcagggtcc | atgtcttagt | acgaaccatc | ctgaagctca | ggagcccgaa | ggttccactg | 240 |
| cctggggaag | gcggc | | | | | 255 |

<210> 1258

<211> 630

<212> DNA

<213> Homo sapien

<400> 1258

| | | | | | | |
|------------|------------|------------|-------------|------------|-------------|-----|
| aaaactaaaa | gcatactgac | tgaactccag | ctcagtcttc | ccattttata | atgaggactc | 60 |
| tgaagtttat | agaggtcaag | gacttgacca | aagctttaga | tatgtagtgt | ctgtgccctt | 120 |
| ttcctctaag | tttctcctag | agaatgtggg | ggctcaggaa | cagagaaaat | aaggtgcaaa | 180 |
| aagtagaaat | gggtggtggt | tctcaaagtg | tggtccatct | gcatactagt | gactgggggtg | 240 |
| cttgttaaaa | tgcagattgc | tgggccttat | cccaatctga | ccaaatcatc | tcaggatcta | 300 |
| ccttttgaac | aaacttgcc | aggtcaaatt | cactcttggtg | gaagtttaag | tacttcagaa | 360 |
| acaagacagc | cacagaaggt | gcacctgcta | atttggtggc | ttccagtgcc | tcatctgtaa | 420 |
| cttctggtga | aatcctgaga | tgtcttactt | tacattgttt | acatcccata | acattccaac | 480 |
| atttagaaat | tactcagagc | ttatttttct | tacttgttta | gcactaaatg | aaaatagctc | 540 |
| cctgaagtta | aggagtttat | atacagtaat | tcatagcaagt | gtgtaaatta | aacagatgac | 600 |
| tttccccctt | aatatcta | gcacagcaag | | | | 630 |

<210> 1259

<211> 159

<212> DNA

<213> Homo sapien

<400> 1259
 aaaattttaca gataaaggca gttcaatact gccactgaga agtacatctc ttaacatata 60
 caacttttcag gccacagttt tgaaggctcg aagtattaag ttggtttgat gaattagtcg 120
 gttggcactt acgaacacat ttattgcctt gccatcttt 159

<210> 1260
 <211> 115
 <212> DNA
 <213> Homo sapien

<400> 1260
 aaaaatacta taattttcaaa acttccaaat ttcaacagat gccagtgttc tctccttttt 60
 tcatatggga aaattttcttt caaaattatt tgacgcttgg acaaaaattc cacag 115

<210> 1261
 <211> 280
 <212> DNA
 <213> Homo sapien

<400> 1261
 aaaatattgt ttatctttat ttattttgtg gtaatatagt aagttttttt agaagacaat 60
 tttcataact tgataaatta tagttttgtt tgtagaaaaa gttgctctta aaagatgtaa 120
 atagatgaca aacgatgtaa ataattttgt aagaggcctc aaaatgttta tacgtggaaa 180
 cacacctaca tgaaaagcag aaatcggttg ctgttttget tctttttccc tcttattttt 240
 gtattgtggt catttcctat gcaaataatg gagcaaacag 280

<210> 1262
 <211> 144
 <212> DNA
 <213> Homo sapien

<400> 1262
 aaattatttg atgagttcca cttgtatcat ggctaccgcg aggagaagag gagtttgtaa 60
 actgggccta tgtagtagcc tcattttacca tcgwtgtgat tactgaccac atatgcttgt 120
 cactgggaaa gaagcctggt tcag 144

<210> 1263
 <211> 487
 <212> DNA
 <213> Homo sapien

<400> 1263
 aaacatcttg ataatttggt gttgagagct gttcattcta aaatgtaatg aaattcagtc 60
 tagttctgct gataaagatc atcagttttg aaaggttact gattttcctc ttccctctta 120
 gttttttacc caatatatgg agaagagtaa tggccaatct taacattttg ttttaattgt 180
 ttaataaagc tgctgggcag tgggtgcagca ttcctaccta gtgtcataaa agcaaaaatac 240
 ttacatagct ttcttaaaat ataggaatga cattacattt ttaggagaaa gtaagttgct 300
 ttgcaccgcc tacttaattc ttttccatat attgtgatac aaacttttga atatggaatc 360
 ttactatttg aatagaaatg tgtatgtata atatacatc atacataagc atatatgtgt 420
 gtgtgtgtgt gtatatatat atatatgcat gctgtgaaac ttgactacac aacataaatc 480
 acttttt 487

<210> 1264
 <211> 250

<212> DNA

<213> Homo sapien

<400> 1264

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgcttcaac | agagtggcag | caaccaagct | ggagtccaag | ccccctgata | aaaggcagcc | 60 |
| aatccttctg | tctgtcatca | aacgtttctt | tacagcatta | ttaaaaagga | tcctgaggtt | 120 |
| gttcttcaca | gtttctatct | caaaacctgg | aaagagtttc | tccacattgt | catagagggc | 180 |
| gtgcaggggt | tcaccccgac | agtgatgata | tttaaccatt | tccacggatg | caactttgcc | 240 |
| atttggcttt | | | | | | 250 |

<210> 1265

<211> 394

<212> DNA

<213> Homo sapien

<400> 1265

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| aaatatttgt | tccaaccttt | ttcgttgggtg | gcatttatgg | ctttggagca | ctgtcaggcc | 60 |
| catgttcatt | accgtgagct | cctgtgcac | tcctaatttc | caaactagcc | tggaacacgc | 120 |
| ctccattgac | catgattggt | tcattggcct | gtgcatggaa | catcatatgt | tcaggagat | 180 |
| aaagaactct | gatagtggca | cctgggtaaa | aagtacaatc | cattatatct | ggatatcaag | 240 |
| atcttttgca | gttgaagaga | ggtattgcca | cagagaaaat | tataggagca | gaagaaagtc | 300 |
| aatgaaagtc | aatgatgaca | ctccattagg | aaccagaaag | atggtattta | tttatacata | 360 |
| taataggtgt | aagagattag | aggaagcctg | tcac | | | 394 |

<210> 1266

<211> 229

<212> DNA

<213> Homo sapien

<400> 1266

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccacagttgt | atcatatagc | atctctaaca | tttcatctag | gattatctag | tatagatctt | 60 |
| actatatttg | gggctatggt | gtatacaatg | ttaacaagaa | catatcttct | ctgcatatat | 120 |
| gtgtgaatta | taaagaaaag | catgagaatg | actctaagtt | caacaaacat | gggtgaatct | 180 |
| ctatgtgtct | ccagtgtcct | ggatgggtct | cccagcaagc | cattcctcc | | 229 |

<210> 1267

<211> 722

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(722)

<223> n = A,T,C or G

<400> 1267

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| aaatcttata | aactttccaa | attttcatac | taaaatatat | tattgtatta | atacaaaacta | 60 |
| cagtattata | cactacactg | tgtaataaat | aaagaaatat | aaaaataaga | cacataaata | 120 |
| taaaagtttt | ctaaaactaa | aagtacatat | gtcagtaaga | agggatttaa | tactgccagg | 180 |
| tttgaagaca | tacagtacaa | aaatgttgca | cagatctata | aactaaaaga | aataaaaataa | 240 |
| tactgatagg | taaaaatcag | ctaattgtgt | taataaattg | gtcccataat | aactaacatt | 300 |
| tggaacacag | tatgagccaa | ataacaatag | catgtccatg | tctgaaatgc | aagtacatgg | 360 |
| ataaagcaga | ttagaaaatt | tccctttcgt | ttctgtagag | aaattctgaa | aatcaatcaa | 420 |
| cataaaatca | ataccgagga | attgaaggat | gaaatgtccc | agtgtttcag | tttctctgac | 480 |

agagtcagtg gttttaagtt ttatttgga attttgatac aagagacaaa tcaacaaatg 540
 ctagttattg taggccacac attggatgaa ggcgggtag agccttgaaa atactgagaa 600
 atggcactta cagcacacag gtcttgctta agggcaaagg agatacaaag cttcatgnca 660
 tatccttcat atggtaccac atattcaaac accatcccaa cactgatctg atgattttgc 720
 tg 722

<210> 1268

<211> 407

<212> DNA

<213> Homo sapien

<400> 1268

gatgacacaa gcagctaata accattttctg gggtttctgcc taaccccccta attgtctgtt 60
 aaagccaatt ctctgggtgt ccagtgagt ggtggctttt tttctttcca cattggcaca 120
 ttcacttctc ccactcttg catgtaagaa ataagcattt acataattgg aaaaatctgg 180
 atttctgatg ccaaagggtt aaagcttctt ggatttcatt tcattgatat acagccacta 240
 ttttattttt gatcagtggt ctttgggcca ctgttcaggg tactgaccat cagtgtcagc 300
 attaggggtt tggtttttgt ttcttttggg tattttcttt ttggcacatg tgaatcttgt 360
 tttgtgtaaa atgaaattac tttctcttgt tctctgatga tgggttt 407

<210> 1269

<211> 675

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(675)

<223> n = A,T,C or G

<400> 1269

ctgaaaaaga gtgatectca atatcctaac taactgggtcc tcaactcaag cagagtttct 60
 tcaactctggc actgtgatca tgaaacttag tagaggggat tgtgtgtatt ttatacaaat 120
 ttaataacaat gtcttacatt gataaaattc ttaaagagca aaactgcatt ttatttctgc 180
 atccacattc caatcatatt agaactaaga tatttatcta tgaagatata aatgggtgcag 240
 agagactttc atctgtggat tgcgttgttt cttaggggtc cttagcactga tgccctgcaca 300
 agcatgtgat atgtgaaata aaatggattc ttctatagct aaatgagttc cctctgggga 360
 gaggttctggt actgcaatca caatgccaga tgggtgtttat gggctatttg tgtaagtaag 420
 tggtaagatg ctatgaagta agtgtgtttg ttttcatctt atggaaaactc ttgatgcatg 480
 tgcttttcta tgggaataaat tttgggtgcaa tatgatgtca ttcaactttg cattgaattg 540
 aaattttggg tggatttata tgtattatac cctgtcacgc ttctagttgc ttcaaccatt 600
 tataccattt tgnacatatt tttacttgna aatatttacc tgncccggcc ggccgtcgaa 660
 agggcgaaat tcaac 675

<210> 1270

<211> 268

<212> DNA

<213> Homo sapien

<400> 1270

ccatcctggg cggagctaaa gttgcagaca agatccagct catcaataat atgctggaca 60
 aagtcaatga gatgattatt ggtgggtggaa tggtttttac cttccttaag gtgctcaaca 120
 acatggagat tggcacttct ctgtttgatg aagagggagc caagattgtc aaagacctaa 180
 tgtccaaagc tgagaagaat ggtgtgaaga ttaccttgcc tgttgacttt gtcactgctg 240

acaagtttga tgagaatgcc aagactgg

268

<210> 1271

<211> 307

<212> DNA

<213> Homo sapien

<400> 1271

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| cctactcttc | tccgtccatt | gtactatctg | cccgtggtgg | ggatggcagt | aggatcatat | 60 |
| ttgatgactt | ccgagaagca | tattattggc | ttcgtcataa | tactccagag | gatgcgaagg | 120 |
| tcatgtcctg | gtgggattat | ggctatcaga | ttacagctat | ggcaaaccga | acaatttttag | 180 |
| tggacaataa | cacatggaat | aatacccata | tttctcgagt | agggcaggca | atggcgtcca | 240 |
| cagaggaaaa | agcctatgag | atcatgaggg | agctcgatgt | cagctatgtg | ctggtcattt | 300 |
| ttggagg | | | | | | 307 |

<210> 1272

<211> 798

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(798)

<223> n = A,T,C or G

<400> 1272

| | | | | | | |
|-------------|-------------|------------|------------|-------------|------------|-----|
| ccattgctag | aaattgaatc | acaaataata | gctaataatt | tttcattttt | caaaaaagat | 60 |
| catttgata | gcagctatgt | ataaaatgga | aaataaaaaa | ttattctatt | ttgcatgaat | 120 |
| agttcagact | ttcccatacc | acagccaagc | agtaactaaa | attaggatct | taattttcaa | 180 |
| tgataaaaagg | tctaaggttc | atttaattat | gctcctttta | cactgtcttt | ctagattttt | 240 |
| caccagat | tttcaaaatt | tgggaatgta | aacaattgat | atattttattg | tatggtggct | 300 |
| agcagttcat | ccttctgcaa | aatatgcatt | cagagaaatg | tgaagcttgt | tttaatgaag | 360 |
| acttaaacca | tttgtgtcat | tttgtttttc | atattcaaat | acaccaaatt | aaaattctga | 420 |
| acctatattt | ttcatcatta | acttccta | ataccagaac | atataccttt | ttcatgtaaa | 480 |
| gttggaatg | ggatattggca | gttttatttt | tgaaaaatat | gtaacatgac | tttaatat | 540 |
| ttatagtttt | cagaattaga | aacataggaa | gggaaaatgt | tttaattaga | taagtcaact | 600 |
| ttttatgggc | tgnagtggng | actataatag | caaattataa | agcattatta | aatgggtata | 660 |
| ataattttta | tattacctca | ttatgaatta | actaaaataa | agnngagtga | tattttta | 720 |
| gggtgntcat | actggagctc | ctgagatata | tgatttgcta | ttgactcact | ggntgattga | 780 |
| ataatatatt | actcgcgg | | | | | 798 |

<210> 1273

<211> 664

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(664)

<223> n = A,T,C or G

<400> 1273

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaaatatacc | ttttcacagg | tagcaagaaa | tagtacatgt | aataagtctt | tatgactgga | 60 |
| atgatccaga | aatatcacia | agcatgagta | aacacatata | taaaagtagc | tcatcatttc | 120 |

```
<210> 1274
<211> 153
<212> DNA
<213> Homo sapien
```

```
<210> 1275
<211> 504
<212> DNA
<213> Homo sapien
```

```
<210> 1276
<211> 533
<212> DNA
<213> Homo sapien
```

<210> 1277

<211> 78
 <212> DNA
 <213> Homo sapien

<400> 1277
 ccacaggaag ttgcaaaaat tagatggact ctgtgtagct agccactctt gagtgtcagg 60
 tctgcatatg tgagtttt 78

<210> 1278
 <211> 560
 <212> DNA
 <213> Homo sapien

<400> 1278
 aaatatctaa aacaatggcc cactgaagaa aggaacaatt aactcttta ttaattcctt 60
 aggataagta ccagaaaatt taacagctag ggcagacttc taatacaata ccgaaagtcc 120
 ttccaaaaac caagtgggtg ccaacttatg tcccttagca ttataacatt cttgagccaa 180
 tagtgtaaaa atacgctgac aatttttatg gcaaacatta ctcaagggtat cttactttcc 240
 acttattact aaagtaatta acccctaaat agatgctcct caacagtggg actacatcct 300
 ggtaaaccta tcataagttg aaactatcaa gttgaaatgc atttagtacc cggataaacc 360
 tatcataaag ttgaaaattt gttaaattgaa ccagtgtaaa tcagaggcca tcttacttca 420
 tactcatgaa gcaactatag tgggatattt ttcaacttac gagatagcct aggcttggtg 480
 aaacactgtc ctaatttact ggctctctgg taattaagtc ataaatggtc aaacatcaaa 540
 ttctagaaaa gcatatattt 560

<210> 1279
 <211> 580
 <212> DNA
 <213> Homo sapien

<400> 1279
 aaaggagatt gtttcaaaaat atttttgcaa attgagataa ggacagaaag attgagaaac 60
 attgtatatt ttgcaaaaac aagatgtttg tagctgtttc agagagagta cggatatatt 120
 atggtaattt tatccactag caaatcttga tttagtttga tagtggtgtg aattttatatt 180
 tgaaggataa gaccatggga aaattgtggt aaagactgtt tgtacccttc atgaaataat 240
 tctgaagttg ccacatggtt tactaatctt ctgtgaaatg catagatatg cgcattgtca 300
 actttttatt gtggtcttat aattaaatgt aaaattgaaa attcatttgc tgtttcaaa 360
 tgtgatattc ttcacaatag cttttttata gtcagtaatt cagaataatc aagttcatat 420
 ggataaatgc atttttatatt cctattttct tagggagtgc tacaaatgtt tgtcacttaa 480
 atttcaagtt tctgttttaa tagttaactg actatagatt gttttctatg ccatgtatgt 540
 gccacttctg agagtagtaa atgactcttt gctacatttt 580

<210> 1280
 <211> 307
 <212> DNA
 <213> Homo sapien

<400> 1280
 aaacacatac gaagaaatca actgtgatta tgaagtggca gccagctaaa tatgtcttgt 60
 atttgtctct ttcctttttt tgccctaactc atcctttact tccattcctg cttccatggt 120
 aatgcaggct caaataaatt actaggatac aagattactt caagcctctt ttctgtggaa 180
 ctcataatat gataagcatt tgttacaaga ttgctgttag ttgtttaggg gataaattat 240
 attagggaaa gaaagtcttt ctttagtttg tttaaatttc tattataatt ggttactaaa 300
 tttattt 307

```
<220>  
<221> misc_feature  
<222> (1)...(745)
```

[illegible]

| | | | | | | | | | |
|-------------|------------|------------|------------|------------|------------|--|--|-----|--|
| <400> | 1285 | | | | | | | | |
| cgacggtatc | gataagcttg | atatcgaatt | cctgcagccc | gggggatcca | ctagtтата | | | 60 | |
| atagtaatca | attacggggt | cattagttca | tagcccatat | atggagttcc | gcgttacata | | | 120 | |
| aacttacgcta | aatggccgcc | accgcggttg | agcttcaggc | tttgttccct | ttagtgaggg | | | 180 | |
| ttaaattqgc | | | | | | | | 190 | |

```

<400> 1286
ctgcatcttt ctacaattct accagcaata tatgagggtt acaattttctc yccatctttg      60
tgaacgcttg ttagagtctg tctctttttc ttccattctg tgggttggct ttttactttc      120
taaatggtag aaccttcaaa gcacaaaggt ttt                                     153

```

| <400> 1287 | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| aaaaacacaa | aacactagaa | cagttgctat | gaaattactg | ataatgatcc | ctttaataaa | 60 |
| ctgcaattaa | ccactaatat | agaaaattcaa | tttaagcaag | aagttttata | tattatactt | 120 |
| tacagaaaaa | aataattttg | aaaaagtaat | gmcaaacaga | gatcaaacat | ttagggcatt | 180 |
| agttactgca | ttctcttttt | agaatataca | ttaagtaaca | ctagtaaaat | tt | 232 |

```
<210> 1288
<211> 90
<212> DNA
<213> Homo sapien
```

<400> 1288
 aaacttagtg actatntagt tcaattgytc atccattttt tatttgcttt tataattgcc 60
 tccttgtttt ggtatattgt aaaataattt 90

<210> 1289

<211> 670

<212> DNA

<213> Homo sapien

<400> 1289
 aaatcacaaa gtaaggcacc attggattaa acattttctcc tggctttttac taagtaaaat 60
 gcatagtga ataaatactg aacactgagt tttaatactg taatacattt caatataaaa 120
 taagaggtga atgttaaaat actgtattac atgttgaata ctttatctg aaaatgttat 180
 aaaaaaacac acatgtaagc tctgatttca gggaagaaaa attcattttt gtaattttcc 240
 atagttaaag attttaccac agaacttatt catagtttta gatgcaatta ggttgcaaac 300
 tttcaaagaa aggggtgtagg tgtattaatg aaacagtcac ttaaacta ctttctaaaa 360
 caatctattc tggatgaatg gcaactttga gctatcacc tgtttcagat ttagaacggt 420
 acctgccaa gttcagatat caaaggaatt gtccaattct tactaccct tataaaattc 480
 agactcactt tctctgagtc agacttttct ccgtcatatt ttctaggaag ggcaaattcc 540
 atcttttggt aaatgggtca ttaggcttta tcatagggat gtttttctact gttgaaatca 600
 gataaaagaa tcccaaataa atgatgctgc taaattacca aactgctaga gattaaaaaa 660
 attttttttt 670

<210> 1290

<211> 352

<212> DNA

<213> Homo sapien

<400> 1290
 aaacaatgct acaccattt ttggcaaagt gctgtattgt tcagtctgtg tacaaaactg 60
 accatctatg aaccaatcag tataaaaaat ttctataaaa acaaaattta gacagtggct 120
 caagaaaaca agctgccatt tatgcataga ttgatgtaca gtaacctaac caaatgtccc 180
 ttttgaattt tcaagttact gaaaaaaaaat gtgtcgagaa acacattaag aaggcacatg 240
 tacagtctac aatactcttc agtctcccta actcatgccc tgcccctata aaggaaatat 300
 gttcacaaat ttacttgaga aaaaaaaaaa aagccactta aaaaaaaaaa aa 352

<210> 1291

<211> 99

<212> DNA

<213> Homo sapien

<400> 1291
 aaaaattatt taaggtaatg gtgttacgaa tggtttaaaa atgtctggtg acttgcttat 60
 ttttaagtga tcaccattaa gtcagaaaaa tgtattttt 99

<210> 1292

<211> 295

<212> DNA

<213> Homo sapien

<400> 1292
 aaatatacct ttattttctca aactcaaagc tttatcaagt tctaacacat tttgcattga 60
 caagtgattt tatctgcac aagtaagggt agtgaccacc acgaaagagg aatccccaga 120
 cctcctaggc actaagaaat atttcaaagg ctatgcaaat atagaacaaa aagcttttcaa 180

```
<210> 1293
<211> 256
<212> DNA
<213> Homo sapien
```

```
<210> 1294
<211> 90
<212> DNA
<213> Homo sapien
```

```
<210> 1295
<211> 519
<212> DNA
<213> Homo sapien
```

```
<210> 1296
<211> 419
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|------------|------------|------------|-------------|------------|------------|--|-----|
| <400> | 1296 | | | | | | |
| aaagcaaac | gcagaaacca | gaagcttctg | accctctaac | atgtattact | gtccaaccca | | 60 |
| ccatgagaag | tatgttcact | tggtgacaac | aaagagactc | cgtatcatat | gtatgttaat | | 120 |
| gaccagattg | ttcatatggg | atTTTTctta | acagattatc | agggtgagaa | tgattctttt | | 180 |
| tctccaaggg | caagaaaaag | ctggctaaat | gctagttaat | taaatccatt | ctcaattttg | | 240 |
| aactgtagag | aagaacctga | cttgaatgag | atTTTTctaaa | ggaagacatt | tcttgctcaa | | 300 |
| cctcaggtat | aattagatta | taaggaatct | cacgtccaga | atTTTatctg | ctgattgtta | | 360 |
| gtatggtagg | taattggcct | taggacacta | tttctactag | aaccctttac | attatttttt | | 419 |

<223> n = A,T,C or G

<400> 1300

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccntngaatt | gtgtgcatag | ggaagcactc | acccaatgag | actttctcca | atgtggactc | 60 |
| tgtgtgtcag | ggaatgaatg | tagaaaaatt | cactttggag | ggttatcakc | tcaactagta | 120 |
| agaagcatta | atattattaa | agtgaagaaa | ctgcagagaa | aattacagaa | caaaactgta | 180 |
| gg | | | | | | 182 |

<210> 1301

<211> 312

<212> DNA

<213> Homo sapien

<400> 1301

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaagttttta | tctctgctga | ggcttcacat | ctgtttgctc | aattttatct | ttatttcaat | 60 |
| ccttgagcat | gtttataata | tagtagtata | cccttattgt | ggctttactt | tcctcacttt | 120 |
| cagtcaccca | cagtcaaaaa | atatgaaata | taaaactcca | gaagtaaaca | gtttataaat | 180 |
| tttaagtcac | actttgttct | gaggaatgtg | atgcaacctc | ccgccattct | gctgtatcca | 240 |
| gttcaggatg | tgacataccc | ctttgctcag | cagatacaca | attcctgctt | cctgctcatt | 300 |
| agacatttgc | ag | | | | | 312 |

<210> 1302

<211> 109

<212> DNA

<213> Homo sapien

<400> 1302

| | | | | | | |
|-------------|------------|------------|-------------|------------|------------|-----|
| attccttagat | tatatgtgtc | catccttgca | gctttctgag | agtaatttta | tttgttgtct | 60 |
| tctgaaatgt | acatgtatac | atgtacctac | tgagtgtctat | gtgattttt | | 109 |

<210> 1303

<211> 330

<212> DNA

<213> Homo sapien

<400> 1303

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccagagttac | ttggatcagc | atttaggaaa | gtaaaatata | gtggaagtaa | aactgactca | 60 |
| tccaactaga | cattctacag | aaagaaaaat | gcattattga | cgaactggct | acagtaccat | 120 |
| gctctcagc | cagcccgtgt | gtataatatg | aagaccaa | gatagaactg | tactgttttc | 180 |
| tgggccagtg | agccagaaat | tgattaaggc | tttcttttgt | aggtaaactc | agagtttata | 240 |
| cagtgtacat | gtacatagta | aagtattttt | gattaacaat | gtattttta | aacatatcta | 300 |
| aagtcacat | gaactggctt | gtacattttt | | | | 330 |

<210> 1304

<211> 170

<212> DNA

<213> Homo sapien

<400> 1304

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ccactgtagt | ctgcatatcc | ctgtccatat | ccatagttcc | catagttata | cccagtataa | 60 |
| tcatatccgc | catagccact | atagttttga | tcaccaccat | aggcactatt | gtaattttcca | 120 |
| tatccttgat | cataatagtt | attaaatcct | tggttccagt | tttggccctg | | 170 |

<210> 1305

<400> 1308
 ctgtcttttg gaggacgtac gtaataaggt tttaatttag taaaccaatc ctatgcatag 60
 tttcagcaact agccaaacct caccaactcc tagttctaga aaaacaggca cttggcagcc 120
 ttgtgatgtc atacagagaa gtcacaggca gtacctgagg gtctgtaggt tgcacacttt 180
 ggtaccagat aacttttttt ttctttataa gaaagcctga gtactccaca ctgcacaata 240
 actcctccca gggttttaac tttgttttat tttcaaaacc aggtccaatg agcttttctga 300
 gcag 304

<210> 1309
 <211> 289
 <212> DNA
 <213> Homo sapien

<400> 1309
 gggatttcca attaacagta ttaccagata aatattcttg gtccaagcag aaaatatcaa 60
 caaaaagagc cttcttctcc tgtaaatctt aaatgcctac atcactcttt atgatacatg 120
 gatcatctta tgtggatact taaatttttc atgtctgctt cttttgcctc tcccaactat 180
 actatgagga aattcggaac aaagacattt ttgtaatat tcttatctcc ttcacaccta 240
 gtatagagct gattttacaa aggcatttaa gagatatttg aattgattt 289

<210> 1310
 <211> 534
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(534)
 <223> n = A,T,C or G

<400> 1310
 tgctttgcat tttctgatgt attacatgac tgtttctttt gtaaagagaa tcaactaggt 60
 atttaagact gataatttta caatttatat gcttcacata gcatgtcaac ttttgactaa 120
 gaattttggt ttactttttt aacatgtggt aaacagagaa aggggccatg aaggaaagtg 180
 tatgagttgc atttgtaaaa atgagacttt ttcagtggaa ctctaaacct tgtgatgact 240
 actaacaat gtaaaattat gaggatgtaa gaaaacattg ctttgtggtt atcactttaa 300
 gytttgacac ctagattata gtcttagtaa tagcatccac tggaaaagggt gaaaatgttt 360
 tattcagcat ttaacttaca tttgtacttt agagtatttt tgtataaaat ccatagattt 420
 attttacatt tagagtattt acactattga taaagtttgt aaataatttt ctaagacagn 480
 ttttatatan gctacagggt gccctgattt tcttattgaa tttgggttaga ctag 534

<210> 1311
 <211> 114
 <212> DNA
 <213> Homo sapien

<400> 1311
 aaaatttgta ggagttgtag actacctaaa tttttaagtt atggyatttg gtcataaggtt 60
 gactgggtag gtaaagaagg aaacagacaa gaaaatggct tcttgagggt gcag 114

<210> 1312
 <211> 95
 <212> DNA

<213> Homo sapien

<400> 1312

```
gggcgggtaa aggtaggccg cgagagcgag gttaggagag gataggaggc cgcagtactg      60
ctcacacgct ccgctcttct cccactctcg actct                                     95
```

<210> 1313

<211> 519

<212> DNA

<213> Homo sapien

<400> 1313

```
aaatgataca gtatttttagg tatgatttaa gactatgatt tacctataca ttatatatat      60
tttataaaga tactaaacca gcataccctt actctgccag agtagtgaag ctaattaaac      120
acgttttggt tctgaataaa ttgaactaaa tccaaactat ttcctaaaat cacaggacat      180
taaggaccaa tagcatctgt gccagagatg tactgttatt agctgggaag accaattcta      240
acagcaaata acagtctgag actcctcata cctcagtggt tagaagcatg tctctcttga      300
gctacagtag aggggaagggt attgttgtgt agtcaagtca ccatgctgaa tgtacactga      360
ttcctttatg atgactgctt aactccccac tgctgtccc agagaggctt tccaatgtag      420
ctcagtaatt cctgttactt tacagacagg aaagttccag aaactttaag aacaaactct      480
gaaagaccta tgagcaaata ggctgaatac tttttttttt                               519
```

<210> 1314

<211> 518

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(518)

<223> n = A,T,C or G

<400> 1314

```
ccatgggtggg tgaagacgct gatctgccct gtcacctggg gttttttatg agtgcagaga      60
ccaggagct gaggaacccc gagytcacgc ctaaggcagg tggngaacgt gtatgcagat      120
ggaaaggaag tggaagacag gcagagtga ccttatcgag ggagaacttc gattctgcgg      180
gatggcatca ctgcaggga ggctgctctc cgaatacaca acgtcacagc ctctgacagt      240
ggaaagnact tgtgttattt ccaagatggn gacttctacg aaaaagccct ggtggagctg      300
aaggttgcag gtgagcctcc aggttttgnt ctgagaacac ttctctgtag gatctanagc      360
agatgcagag tccctcttcc aaaagtactg cagacactcc tggctgctca ctagcaatng      420
tctgcactgc ctcccaactn agcttctctg caacccttaa gaaagacaca ttctttcttt      480
agaaagaatt cctgctgnac cttacatgcc gaagtaaa                               518
```

<210> 1315

<211> 360

<212> DNA

<213> Homo sapien

<400> 1315

```
tctgtgcac caatttatta tagwtttgta agtaacaata tgtaatcaaa cttctagggt      60
acttgagagt ggaacctcct atatcattat ttagcaccgt ttgtgacagt aaccatttca      120
gtgtattggt tattatacca cttatatcaa cttatttttc accagkataa watcttratt      180
tytacgacct atcattctga atcaagmaca ctgtatgttc agtaggttga actatgaaca      240
ctgtcatcaa tgttcagttc aaaagcctga aagtttagat ctagaagctg gtaaaaatga      300
```


<400> 1319
aaatttagtg tctcatttgg aaataaactc tgggcctatt agttgttgag tattttttttt 60
ttttactacc taaaaaaaga tttgttaaga gctgaattac aacttagcat tacataatat 120
aaaacactgt aatgtgtatt t 141

<210> 1320
<211> 497
<212> DNA
<213> Homo sapien

<400> 1320
aaattcagtc ctaagaaaga ggagtgcttg tcccctaagg gtgtttaatg gcaaggcagc 60
cctgtctgaa ggacacttcc tgcctaaggg agagtggtat ttgcagacta gaattctagt 120
gctgctgaag atgaatcaat gggaaatact actcctgtaa ttcctacctc cctgcaacca 180
actacaacca agctctctgc atctactccc aagtatgggg ttcaagagag taatgggttt 240
catatttctt atcaccacag taagtctcta ctaggcaaaa tgagagggca gtgtttcctt 300
tttggtactt attactgcta agtatttccc agcacatgaa accttatttt ttcccaaagc 360
cagaaccaga tgagtaaagg agtaagaacc ttgcctgaac atccttcctt cccacccatc 420
gctgtgtgtt agttcccaac atcgaatgtg tacaacttaa gttggtcctt tacactcagg 480
ctttcactat ttccttt 497

<210> 1321
<211> 344
<212> DNA
<213> Homo sapien

<400> 1321
ctgtccaatg acaacaggac cctcactcta ctcaagtgtca caaggaatga tgtaggaccc 60
tatgagtgtg gaatccagaa cgaattaagt gttgaccaca ggcacccagt catcctgaat 120
gtcctctatg gccagacga cccaccatt tccccctcat acacctatta ccgtccaggg 180
gtgaacctca gcctctcctg ccatgcagcc tctaaccacac ctgcacagta ttcttggtctg 240
attgatggga acatccagca acacacacaa gagctcttta tctccaacat cactgagaag 300
aacagcggac tctatacctg ccaggccaat aactcagcca gtgg 344

<210> 1322
<211> 110
<212> DNA
<213> Homo sapien

<400> 1322
ccaccacata gccagccagg aatcccttga ggaacgggga ggacaacagc gagccacctt 60
ggcccactcc actgttgact tcgtcttcta cacgccgctg caggctttcc 110

<210> 1323
<211> 359
<212> DNA
<213> Homo sapien

<400> 1323
ccacgtgctt ggccctgggct ggcgtctcct gctgtgagct ggctgaggag gacttctctg 60
cggctctccc cttagatccg cgctatcgtg aggtccacta tgcctgctg gatccttctt 120
gcagtggctc ggggtgagatg gtgagaaggc gtggctgagg gactcagagg tccacagcag 180
cttagacctg gagtcatctg ttttggtcct agttctgaca ctttaatggg cttggggacct 240
tgagagcaaaa gttctcctct gtgaagcgag gatttcagga gcgaggattt caggactgag 300

gcagcctgtg aagctgtgta accgagacac gcttttcctt aggtatgccg agcagacag 359

<210> 1324

<211> 258

<212> DNA

<213> Homo sapien

<400> 1324

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| caatcacaca | accacaaaaa | agatactgtg | tgctctcact | ttccaaaatt | ctgcctggtc | 60 |
| tmctcctgag | gaaagyagtg | atatggtagc | tggtgtggat | cccctaaagg | aattataaga | 120 |
| tggartgyga | rgaacattat | cttagactat | aakactgkct | gcatrcrgat | atgktstcra | 180 |
| agattattcc | tgctgcraat | aaagakmttg | skaaagagca | rtatasagct | atcacagtct | 240 |
| attgacccam | asatgttt | | | | | 258 |

<210> 1325

<211> 534

<212> DNA

<213> Homo sapien

<400> 1325

| | | | | | | |
|------------|------------|-------------|-------------|------------|-------------|-----|
| ctgtccaatg | gcaacaggac | cctcactcta | ttcaatgtca | caagaaatga | cacagcaagc | 60 |
| tacaaatgtg | aaaccagaa | cccagtgagt | gccaggcgca | gtgattcagt | catcctgaat | 120 |
| gtcctctatg | gcccggatgc | ccccaccatt | tcccctctaa | acacatotta | cagatcaggg | 180 |
| gaaaatctga | acctctcctg | ccacgcagcc | tctaaccac | ctgcacagta | ctcttggttt | 240 |
| gtcaatggga | ctttccagca | atccacccaa | gagctcttta | tccccaacat | caactgtgaat | 300 |
| aatagtggat | cctatacgtg | ccaagcccat | aactcagaca | ctggcctcaa | taggaccaca | 360 |
| gtcacgacga | tcacagtcta | tgacagagcca | cccaaaccct | tcacaccag | caacaactcc | 420 |
| aaccccgtag | aggatgagga | tgctgtagcc | ttaacctgtg | aacctgagat | tcagaacaca | 480 |
| acctacctgt | ggtgggtaaa | taatcagagc | ctcccgggtca | gtcccaggct | gcag | 534 |

<210> 1326

<211> 177

<212> DNA

<213> Homo sapien

<400> 1326

| | | | | | | |
|------------|------------|------------|-------------|-------------|------------|-----|
| ctgcattatg | tgtgtttaga | acgagaagtt | gtttgtacag | tattttttcta | ttgaccgctt | 60 |
| ccgtcttgcc | tgaaacctgg | gcattctttc | caatagacag | aaaatcagag | agtcaaactc | 120 |
| gatgogcaat | gagttgttct | gagaccagta | atccacgggtg | ctgcaatttg | ggtttttt | 177 |

<210> 1327

<211> 266

<212> DNA

<213> Homo sapien

<400> 1327

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| aaacttgttt | tatctaatac | tgagcactgt | ttttttgtca | agtattttttt | taagaccaca | 60 |
| taattctttt | tgtctgctca | aggaaaggat | agataaataa | ttggcacaca | tttgtttctc | 120 |
| actgaatttt | acagtagtaa | attaatgtta | taatgtacca | catggagatg | agttggtaag | 180 |
| aaatcatcta | gttccagagc | ccagggatta | taaacagtag | gtgaaataga | tttatgactt | 240 |
| acgaaatatg | ttgtgacaat | atattt | | | | 266 |

<210> 1328

<211> 409

<212> DNA
 <213> Homo sapien

<400> 1328
 ctgtccaatg gcaacaggac cctcactcta ttcaatgtca caagaaatga cgcaagagcc 60
 tatgtatgtg gaatccagaa ctcagtgtgt gcaaaccgca gtgaccaggt caccctggat 120
 gtcctctatg ggccggacac ccccatcatt tccccccag actcgtctta cttttcgga 180
 gcgaacctca acctctctg ccaactcgcc tctaaccat ccccgagta ttcttggcg 240
 atcaatggga taccgcagca acacacacaa gttctcttta tcgccaaaat cagccaaat 300
 aataacggga cctatgctg ttttgtctct aacttggcta ctggccgcaa taatcccata 360
 gtcaagagca tcacagtctc tgcattctgga acttctctg gtctctcag 409

<210> 1329
 <211> 136
 <212> DNA
 <213> Homo sapien

<400> 1329
 ccattttcgc acagtccacc ataaaattga aaagattgac cagagacaga tcatggaggg 60
 cttggcaatc tgtactgatg aagccatgga ccagaagaga agtgagtcaa tgaagagagt 120
 ttctcttttc acatgg 136

<210> 1330
 <211> 311
 <212> DNA
 <213> Homo sapien

<400> 1330
 ctgctaacag ccctaacggt gcaacacaag tacaaaactca ggaacctott cgactgccac 60
 gcccttcacc aacagaagga agacagtggc gccaccacaa gtggcagggc acaggggctt 120
 ctgtgacaac aatatgtcct tctagtatac attcattgca aaggctgccc tgaagtttcg 180
 tttttggaaa taactgttat catacatttt gtatgatgtt gcttgtgggc accatgaaga 240
 gagcctggct gttaaaggaca gagggagcta aaccaacaat gcatggccct gcgtgcccac 300
 aagagggagc c 311

<210> 1331
 <211> 613
 <212> DNA
 <213> Homo sapien

<400> 1331
 ctggggccakg agctgtgccc ggtgacctga gccttcataa gcacacacgt ccattcccta 60
 ctaaggccca gacctcctgg tatctgcccc gggctccctc atcccacctc catccggagt 120
 tgcccaagat gcatgtccag cataggcagg attgctcggg ggtgagaagg ttaggtccgg 180
 ctcagactga ataagaagag ataaaatttg ccttaaaact tacctggcag tggctttgct 240
 gcacggctctg aaaccacctg ttcccacctt cttgaccgaa atttccttgt gacacagaga 300
 agggcaaagg tctgagccca gagttgacgg agggagtatt tcagggttca cttcaggggc 360
 tcccaaagcg acaagatcgt tagggagaga ggcccagggt ggggactggg aatttaagga 420
 gagctgggaa cggatccctt aggttcagga agcttctgtg caagctgcga ggatggcttg 480
 ggccgaaggg ttgctctgcc cgccgcgcta gctgtgagct gagcaaagcc ctgggctcac 540
 agcaccacca aagcctgtgg cttcagtcct gcgtctgcac cacacaatca aaaggatcgt 600
 tttgttttgt ttt 613

<210> 1332

<400> 1335
aaattagttg ctataaattc atcaataactt tttttcccta ttatatatttt ggttctatta 60
ggatttactt aactgaatct tataacaatt cgaggtgaac tgtggcaatg aaaaccagaa 120
acagttaatg agatgcttca gctcacagtt tgaagtgtg agaacctag tattttgctg 180
tacggtagtg agctgtacca aaatatgatg gtttaggttt atgtgcaaga ctttgtgttg 240
tagtctagac aaaggggtgg gcaagagaca tgcaaagctg aagccctgct tgaaaagacc 300
cttcaaggaa gtaaaatggc aggggcagag tgcagcttaa catgttgcta tccctgttgt 360
ttttgagttg gttttggaat ggattcaagt tcttacacaa tttattttga atacaagcat 420
aatctaggtg atttgagtta atgaacttct tttcatgatg tagggaaagc tgaatgtata 480
tatttctaag aagaatttgt ttagcagatt acaagttggc aaaatagact gttcacagaa 540
actaggcaaa aattt 555

<210> 1336
<211> 505
<212> DNA
<213> Homo sapien

<400> 1336
cctggaaaga agcccagcaa aaggttccag atgaagaaga aaatgaagag agtgacaacg 60
aaaaggaaac tgaaaaagat gactccgtaa cagattctgg accaaccttc aactatcttc 120
ttgatattgc cctttggtat ttaaccaagg aaaagaaaga tgaactctgc aggctaagaa 180
atgaaaaaga acaagagctg gacacattaa aaagaaagag tccatcagat ttgtggaaag 240
aagacttggc tacattttatt gaagaattgg aggtctgtga agccaaggaa aaacaagatg 300
aacaagtcgg acttctctgg aaagggggga aggcccaagg gaaaaaaaca caaatggctg 360
aagttttgccc ttctccgctg ggtcaaagag tcattccacg aataaccata gaaatgaaag 420
cagaggcaga aargaaaaat aaaaagaaaa ttaagaatga aaatactgaa ggaagccctc 480
aagaagatgg tgtggaacta gaagg 505

<210> 1337
<211> 385
<212> DNA
<213> Homo sapien

<400> 1337
ctggtgctag tcagagctaa tgacagaatt tcagtttaat aaaaagaccc ccaactgagc 60
acaccatctt gaaaaaagta tacttatcaa acagctttca atcagttcaa gagagacacc 120
ttaattgggg agaggaagaa ttgcagagta gtttgaatc atgccaattc cagatcaata 180
actgcatgtc tgttcttttg tagaaatagc ttttgcctta tattaagtaa tcacatatat 240
attctctcta tttggataag gaaaccttcg ctttatttga caatgtataa tgatatactc 300
ttctaattca cctctgtgtc ttcacaataa acatgagtaa aatttagaca agtcatggta 360
aaggtcaata taattattta ttttt 385

<210> 1338
<211> 350
<212> DNA
<213> Homo sapien

<400> 1338
aaaggtgata ttacacaaaa cctcgtcttt tgttcaactt tggatccatt ggcaattcaa 60
tggcctcaat ctcccaaac tcgccaaagt actccctgat cttttcctca gtggcttcag 120
gattcagacc cccaacgaag attttcttca ccgggtcctt cttcatagcc atggcctttt 180
tagggatcaat gacacggcca tccagcctgt gctcctcttg gtctaggacc ttctccacac 240
tggctgcac tttgaacagg ataaacccaa accctcttga ccgtccagtg ttgggatcca 300
tttttattgt acagtcaacg acctctccaa atttagtaaa atagtctttt 350

<210> 1339
 <211> 443
 <212> DNA
 <213> Homo sapien

<400> 1339
 ctgtcctct agtaataagt tccctggggat aatacattaa ccaacattgg ttgaaacata 60
 cctgagtaat catatcagga tgcattgttaa gctgataaaa caataagatc ccaaaatgca 120
 gtagctcaaa aaaagtagaa gttaatttat ctcctggggg acagctctgg ttctcaaatt 180
 ttacaggctc agaatcacct gcagggtctg tgaaagtaca gattgctgcg ctccgcccc 240
 agagtctctg atttagtagg tgtaggctg aaccaagaat ttgcctttct aacaagctcc 300
 caagtgatgc tgatgacttg taggaatgga tttacttcta ggattagact tcagctcact 360
 ctgtttgctg aactctttct aatatttctt aagttggtag actcyctgct ccaggttctc 420
 aacgtgaagg aaggaacccc cag 443

<210> 1340
 <211> 273
 <212> DNA
 <213> Homo sapien

<400> 1340
 cctcaggaac aggtaggggc agcagaatag aatagcatcc atttcccaga gaaagactgc 60
 ctttacatkt cccatgcttt tagcacaaag cagcgtctgg gccactgtta ccagaggtga 120
 gtttatacat ttacaaaatg cttaaaatct ttgggaagca agaggaagct aaacagaagg 180
 tcccatgtta actgaaggca aattcactca acctctctag taagggaccc atgggcctac 240
 agagtgttcc ctctacaatg tgcagagtgg aaa 273

<210> 1341
 <211> 561
 <212> DNA
 <213> Homo sapien

<400> 1341
 ccatgggccc ggtcacgaac aaaacgggccc tggacgcctc gcccctggcc gcagatacct 60
 cctactacca ggggtgttac tcccggccca ttatgaactc ctcttaagaa gacgacggct 120
 tcaggcccgg ctaactctgg caccgccgat cgaggacaag tgagagagca agtgggggtc 180
 gagactttgg ggagacggtg ttgcagagac gcaagggaga agaaatccat aacaccccca 240
 ccccaacacc gccaaagacag cagtcttcyt caccgctgc agccgttccg tcccaaacag 300
 agggccacac agatacccca cgttctatat aaggaggaaa acgggaaaga atataaagtt 360
 aaaaaaaagc ctccggtttc cactactgtg tagactcctg cttcttcaag cacctgcaga 420
 ttctgatttt tttgttgggtg ttgttctcct ccattgctgt tgttgagggt aagtcttact 480
 taataaaaaa aaaaaatttt gtgagtgact cgggtgtaaaa ccatgtagtt ttaacagaac 540
 cagaggggtg tactattgtt t 561

<210> 1342
 <211> 159
 <212> DNA
 <213> Homo sapien

<400> 1342
 aaagatggca aggcaataaa tgtgttcgta agtgccaacc gactaattca tcaaaccaac 60
 ttaatacttc agaccttcaa aactgtggcc tgaaagttgt atatgttaag agatgtactt 120
 ctcagtggca gtattgaact gcctttatct gtaaatttt 159

<210> 1343
 <211> 76
 <212> DNA
 <213> Homo sapien

<400> 1343
 aaaatgtaaa gccaatctat caccaaaaat ggcataaatg taaacacaag ctaattttat 60
 aatccactgc tatttt 76

<210> 1344
 <211> 726
 <212> DNA
 <213> Homo sapien

<400> 1344
 caaaagcagc ctgaatacgc aactcacgcc aagagggcag cagctctcct gacatccatg 60
 taagaaggct aacacctaata ccacacgcag gcatcctgaa ctcagcagct ctgatccaag 120
 gtactgagtg gagacaaaagc actcggaggt ggcaagatgt tcagcaacca agtaagacac 180
 actggcaagg catccccacc aaaggtgaga agcacaaagc aggccttgag aaacaaacag 240
 tcatgccagg tgcagccaga catcctgcta taagccctga ccctagtacc ccgagttcat 300
 caagtgcctt ggttttgtgt ccataaagca cagagggcac tgaccacccc aaaccagaat 360
 cccaaggaat ccttatggat ggcatagggc ctcagaactg ctgcaggatc attttccttt 420
 tcaggtcgtg gctgaacttg ttcacctga agagctcact gtcataaaat gcagagaggt 480
 tgtggatgtt gatctgacga gccttatcca ccaagtcctt mtcagggacc tcaatagtgt 540
 cctgctgggc cccaaagcgg ttgcgctgat atgtcacstg ctctgccact aactgcttca 600
 gtatgaagag caacagctca ttgtgtcac gccggaatga aaggtagcgg gcaaaagtct 660
 tgcgcatgct gcgcatgacg ctgaacttct gtgtgtctat gaagstctcc akmatcayga 720
 gratgg 726

<210> 1345
 <211> 742
 <212> DNA
 <213> Homo sapien

<400> 1345
 ccagagagcc ctgtcctgtg aggggtggta tcacagtggc aggggttcaat tcagaagacc 60
 ttgagggcag gctgatgttt cctgaatggg cccctgggtg ttgcttgctc ctgactctcc 120
 atttccccat ctgagtggat ttggacctaa tagggcactg gagctgggtc gaatcctgac 180
 tggactactt ggcaacttta tgtctgggag caagttactt aacctcccca agcctgtgtc 240
 tgtgaaatgc gggtaaatga atgtagatgt ttggcagcag ctactccttg ttgagctctc 300
 acagtgaact ctccctgcctc tgccctcctt ccccgccctc cctgggtgcct agcgtcaggt 360
 ctageccactt cctcctgggc cctctcctt tttctgtggc tggctgcctg cccgcctggc 420
 gctggacctt tcatgtaacg ggaatcagca tgtatattct ggtctgggtc gtttctacac 480
 ttaattttgt ttccagtagt atttccctgt accggcagag ttcacaaaca catttgaaga 540
 ggctttttct caggattctt aaccttccaa aggaagtccc atggatgggt ttctagaagt 600
 ctataaatgc tctgaaattg tatttttctg tggaaaagca taacttttat ctgcttggtc 660
 gtgtcaaaa aaagatcatg aatggaatga attgcattga attttatgcc attgggggct 720
 taatactaaa aggatatgga ag 742

<210> 1346
 <211> 573
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(573)
 <223> n = A,T,C or G

<400> 1346
 aaatgcattk ttaacttaca gtatitttcaa cttacgatgt gtttatcasg aagtaacccc 60
 atcataagca gaggagcatc tgtattgCGT aatttgactg gcacagttaa ttaggttctg 120
 ttcagtgwtt tccgtcaaca agatgtttat tgtgtgagta aacaagttaa gccctgtgac 180
 aagctgaata agaatagtct ctccctcagca gcttatagta aacaagggtg gtaatcctta 240
 cattagtggc tagactatca aacgaaatat ataacatgta agaacactaa agacagaatt 300
 actgtggcat agagatagtt agaattgctt cagcctaaga gatgaattag gtaatgcaag 360
 gaggtgaata tgttggcctg caatatgaac aaggcagaga gctgggagag taagatgtaa 420
 gttgctaagg agggatgtgt cttgagtttg gaaaccataa agggaaatca taggtaatgc 480
 tagagtcact gatcttangg agccttgaat aacggtgatg actaaggga tctttatatt 540
 gnggggacta ttggaattaa attggccaga att 573

<210> 1347
 <211> 333
 <212> DNA
 <213> Homo sapien

<400> 1347
 cctggtttct ggtggcctct atgaatccca tgtaggggtgc agaccgtact ccatccctcc 60
 ctgtgagcac cacgtcaacg gctcccggcc cccatgcacg ggggagggag atacccccaa 120
 gtgtagcaag atctgtgagc ctggctacag cccgacctac aaacaggaca agcactacgg 180
 atacaattcc tacagcgtct ccaatagcga gaaggacatc atggccgaga tctacaaaaa 240
 cggccccgtg gagggagctt tctctgtgta ttcggacttc ctgctctaca agtcaggagt 300
 gtaccaacac gtcaccggag agatgatggg tgg 333

<210> 1348
 <211> 185
 <212> DNA
 <213> Homo sapien

<400> 1348
 aaaaaagctt gcagcaagaa aatgccagtg tgcaactggg tgactaaaga ccaaagaaaa 60
 acagttaaaa gggacagctt acttgctctc tgtctcaggt ttaacttctc acctgaaatc 120
 tctcatagcc ctaattaaac acaaacaaaa gtctcttcca tagataggct acttctcagc 180
 ttcag 185

<210> 1349
 <211> 171
 <212> DNA
 <213> Homo sapien

<400> 1349
 gcggcagcga ggggctcgga gaggtgctcg gattctcgta gctgtgccgg gacttaacca 60
 ccaccatgtc gagcaaaaaga acaaagacca agaccaagaa gcgccctcag cgtgcaacat 120
 ccaatgtggt tgcctatgtt gaccagtcac agattcagga gttcaaagag g 171

<210> 1350
 <211> 400

<212> DNA

<213> Homo sapien

<400> 1350

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| ttgtcatatc | atatctatgt | cacctgtgta | ttctgagatt | acacacatac | ctgccaatat | 60 |
| acctgggaaa | ggttatttta | tcacagttac | acttgagttc | ttggcaggca | ggactgagga | 120 |
| agagtaattt | gaaagaagtt | ttacatccta | tttagaagaa | atcactagta | tttccttaaa | 180 |
| taacagggtta | caatagaaag | atactgcctg | gaagttatcc | tttcactttg | gttcattttt | 240 |
| agtttttctt | tatgatttac | atagctgttt | aattcatttg | cttatagtac | aatcctgcca | 300 |
| taaagtatta | aagcacaaga | tacctattat | tccttcaaca | tctgcatttt | tcaagtttta | 360 |
| tactctacat | ccacagtaacg | tcagcagttc | ttgaatgttt | | | 400 |

<210> 1351

<211> 309

<212> DNA

<213> Homo sapien

<400> 1351

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccaggaaaag | gcagtcctga | gggagaagac | aggattcagg | gcagtgctcc | gaagctgtgt | 60 |
| gctcacctgg | ttggctcatc | aaacctggca | acctgtggc | ctgtctgccg | gagctgactg | 120 |
| gatccactca | tcaattcttc | gtccccacta | ctaagaactg | gcatgttttg | ctgggtgtgg | 180 |
| ctctgcactt | caggaatgg | cacaacaggg | ggtagccctc | aaaagcactc | ctttttctat | 240 |
| acctcttctc | aaggccatgt | aagttgccc | tctctacctg | gctgtggaca | aaaggttatc | 300 |
| tgctcttgg | | | | | | 309 |

<210> 1352

<211> 268

<212> DNA

<213> Homo sapien

<400> 1352

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ccacttcac | tgtgtgggaa | cgtggtcagg | ccgggtgctg | gtgtttgaca | tcccagcaaa | 60 |
| gggtcccaac | attgtactga | gcgaggagct | ggctggggcac | cagatgccaa | tcacagacat | 120 |
| tgccaccgag | cctgcccagg | gacaggattg | tgtggctgac | atggtgacgg | cagatgactc | 180 |
| aggcttgctg | tgtgtctggc | ggtcagggcc | agaattcaca | ttattgaccc | gcattccagg | 240 |
| atttgagatt | ccgtgcccct | ctgtgcag | | | | 268 |

<210> 1353

<211> 620

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(620)

<223> n = A,T,C or G

<400> 1353

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| cctgagtaat | tattccatca | tagacaaact | tgtgaatata | gtggatgacc | ttgtggagtg | 60 |
| cgtgaaagaa | aactcatcta | aggatctaaa | aaaatcattc | aagagcccag | agcccaggct | 120 |
| ctttactcct | gaagaattct | ttagaatttt | taatagatcc | attgatgcct | tcaaggactt | 180 |
| tgtagtggca | tctgaaacta | gtgatttgtt | ggtttcttca | acattaagtc | ctgagaaaga | 240 |
| ttccagagtc | agtgtcacaa | aaccatttat | gttaccacct | gttgacagcca | gctcccttag | 300 |
| gaatgacagc | agtagcagta | ataggaaggc | caaaaatctc | cctgggagact | ccagcctaca | 360 |

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgggcagcc | atggcattgc | cagcattgtt | ttctcttata | attggctttg | cttttggagc | 420 |
| cttatactgg | aagaagagac | agccaagtct | tacaagggca | gttgaaaata | tacaaattaa | 480 |
| tgaagaggat | aatgagataa | gtatgttgca | agagaaagag | agagagtttc | aagaagtgtg | 540 |
| attgnggctt | gtatcaacac | tgttactttc | gtacattggc | tggaacagt | catgtttgct | 600 |
| ttcataaatg | aagcagcttt | | | | | 620 |

<210> 1354

<211> 398

<212> DNA

<213> Homo sapien

<400> 1354

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| aaaggattat | ttttatgcaa | agtattctgt | ttcagcaagt | gcaaatttta | ttctaagttt | 60 |
| cagagctcta | tatttaattt | aggtcaaagt | ctttccaaaa | agtaattctaa | taaatccatt | 120 |
| ctagaaaaat | atatctaaag | tattgcttta | gaatagttgt | tccactttct | gctgcagtat | 180 |
| tgctttgcc | tcttctgctc | tcagcaaagc | tgatagtcta | tgtcaattaa | ataccctatg | 240 |
| ttatgtaa | agttatttta | tcctgtggtg | catgtttggg | caaatatata | tatagcctga | 300 |
| taaacacatt | ctattaaatc | aaatatgtac | cacagtgtat | gtgtcttttg | caagcttcca | 360 |
| acagggatgt | atcctgtatc | attcattaaa | catagttt | | | 398 |

<210> 1355

<211> 371

<212> DNA

<213> Homo sapien

<400> 1355

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|-------------|-----|
| ctggytcctc | agtgggaact | gagtcattac | ctgctaaagg | gtagaagagg | agagagagag | 60 |
| gccagagcct | ggggatgggg | cagaagggtgc | agcaggaagg | aagggttagag | tgagaaaaat | 120 |
| ttccaaataa | ggggatgatgt | gtgagtgtc | agaggggtgac | tgaggacatc | tccagcattt | 180 |
| ccattgagga | gggaggaagg | aggggccctt | gggttctggg | gcagatgccg | gcaggggtctg | 240 |
| gatgagatgc | ccccaacctc | aacctgtgtc | ctctgaaaac | acttcaccca | gtcacactga | 300 |
| ggagccccctc | caggcccagg | ggccccctcca | ggtaggcgta | tctcagctcc | tctctggaag | 360 |
| gacccccaca | g | | | | | 371 |

<210> 1356

<211> 338

<212> DNA

<213> Homo sapien

<400> 1356

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|-----|
| gcggcgcggg | cgggcggtaaa | atgtcggttc | caggacctta | ccaggcggcc | actgggcctt | 60 |
| cctcagcacc | atccgcacct | ccatcctatg | aagagacagt | ggctgttaac | agttattacc | 120 |
| ccacacctcc | agctcccatg | cctggggccaa | ctacggggct | tgtgacgggg | cctgatggga | 180 |
| agggcatgaa | tcctccttcg | tattataccc | agccagcgcc | catccccaat | aacaatccaa | 240 |
| ttaccgtgca | gacgggtctac | gtgcagcacc | ccatcacctt | tttgaccgcg | cctatccaaa | 300 |
| tgtgttgtcc | ttcctgcaac | aagatgatcg | tgagtcag | | | 338 |

<210> 1357

<211> 159

<212> DNA

<213> Homo sapien

<400> 1357

| | | | | | | |
|------------|------------|------------|------------|------------|------------|----|
| ctgggtgtgt | gcctctggag | tacttccccg | cagctcctca | ttgctcacat | agtaggcaat | 60 |
|------------|------------|------------|------------|------------|------------|----|

ggcgttgctc tcaaacacac agaatccatc atcacccctca aatgctggga ccttgccggc 120
aggaaatttg cggagaaatt caggggtgcg gttggtttg 159

<210> 1358

<211> 306

<212> DNA

<213> Homo sapien

<400> 1358

cctgtcagag tggcactggg agaagttcca ggaaccctga actgtaaggg ttcttcatca 60
gtgccaaacag gatgacatga aatgatgtac tcagaagtgt cctggaatgg ggcccatgag 120
atggttgtct gagagagagc ttcttgtcct gtctttttcc ttccaatcag gggctcgctc 180
ttctgattat tcttcagggc aatgacataa attgtatatt cggttcccgg ttccaggcca 240
gtaatagtag cctctgtgac accagggcgg ggccgagggg ccacttctct gggaggagac 300
ccaggc 306

<210> 1359

<211> 382

<212> DNA

<213> Homo sapien

<400> 1359

agagggagtc cagcccccaa gccttgtgag gcaactgttar gcagataggg aaaagagggg 60
tccttagatc actggttcaa ggagggatct ggtaggggca gcatttcttc tgggctggaa 120
acagaatggg ggtttcaaga tggcagaacc attccattat tggagctata agcccctaga 180
attgctccat ggcctatctc ggtttccctt ggatctcatc tgctcctgaa ctgcacctgt 240
catggcaagt ccctctccgg ccccatctc ccctgagcca atgtgagtca ggtgaacaaa 300
attcattggg tccccaatca tggcccggtc aatccgtctt ctcttcttct ttcttctcca 360
ccatccagac gttcagctac ag 382

<210> 1360

<211> 365

<212> DNA

<213> Homo sapien

<400> 1360

aaaaaacctt tcaaaataaa acttagtaaa atctagaact gkttcttggc ctacttgaga 60
ggaacttcca tattttcaca gccatctccg aaagcagcag ttgctgtaaa ttaactgaga 120
cttggaatg gtgcagactg tcttggtaga gctgttctta tagcacaatt ttatctggaa 180
aataaacttg taaatgcgtg ctgtatatta atacatgtgt gccatattt atttttatta 240
tctcctgccg gtctttgctc aatgggagat gacagaccaa cttctcaacg tgatttcccc 300
atttcattga atgacattta tatgccactt atgaaaaaaaa tactgctgtg aaagaaatgt 360
acttt 365

<210> 1361

<211> 502

<212> DNA

<213> Homo sapien

<400> 1361

gaggtatgga aaaatatcaa caaggaaata ttagatttga actgctgctt cgtaggcaca 60
cagcacattc tccaggatat accatatgtt aggacacaaa acgggtctca ataaattttt 120
aaaagtcaaa atcttatcaa gtatcttctc agaccacaat ggaataaaaac tggaaatcaa 180
taacaagagg aacttctgaa attgaacaga tacacggaaa tcaaactaca tgttcctgaa 240

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tgaccactgt | gtctatgaag | aaattgattt | taaaaattta | aaaattcttt | gaaacaaatg | 300 |
| aaaatagaaa | cacagcatac | aaaaatgtat | agggtacaac | aaaagaagtg | ctatgaggga | 360 |
| catttatctt | aataaacacc | cacatcaata | aggtagaaag | tttttaaaca | aataacctaa | 420 |
| taaacgcac | tcaaggaact | agaaaagcaa | gaacaaatca | aacctaataa | tagaaggaaa | 480 |
| taaatagtaa | agatcagagc | ag | | | | 502 |

<210> 1362

<211> 545

<212> DNA

<213> Homo sapien

<400> 1362

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ctgattggat | gtctaggaat | gactgaaaga | aaccaaaca | gcctgtccac | tgctgctgtg | 60 |
| ggatggagga | ggcgtaagca | gaaacactaa | cagtatactg | acctcttagc | agaaccgctt | 120 |
| ccattctgga | gatcacggct | gctaaatcca | gcacccccac | ttcattttac | ccccagcata | 180 |
| ttgttctgta | gtcttttctt | gaaacatctt | gattgctttt | cctcggcagc | tttcaaaaaa | 240 |
| ccaaataata | atagttatcc | gtcttctact | tcattggaaga | ttgttttggt | gccctgaccc | 300 |
| tttgaagtgc | ccagttcctg | ccatctgaaa | cctcggcctg | atctgatctc | atgttggaat | 360 |
| ctgcctgtct | ttcacacagg | gctgggtctt | gtcctttaca | tgccagtttt | gcttgtgaat | 420 |
| tcttgctttt | ttcctctcat | cagccttaag | tttaggcgtt | tggtgttctc | cagtgatgta | 480 |
| gacagttccc | ttcacaagtc | acagttcttc | ccataaatga | ggccccgtga | cctctgctgg | 540 |
| acttt | | | | | | 545 |

<210> 1363

<211> 286

<212> DNA

<213> Homo sapien

<400> 1363

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| gggagatgca | ggatgtagac | ctcgctgagg | tgaagccttt | ggtggagaaa | ggggagacca | 60 |
| tcaccggcct | cctgcaagag | tttgatgtcc | aggagcagga | catcgagact | ttacatggct | 120 |
| ctgttcacgt | cacgctgtgt | gggactccca | agggaaaccg | gcctgtcatc | ctcacctacc | 180 |
| atgacatcgg | catgaaccac | aaaacctgct | acaacccccct | cttcaactac | gaggacatgc | 240 |
| aggagatcac | ccagcacttt | gccgtctgcc | acgtggacgc | ccctgg | | 286 |

<210> 1364

<211> 503

<212> DNA

<213> Homo sapien

<400> 1364

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| ccatcaggat | catgaaaaca | aacttttggtg | aatgtgagca | actgcgccag | acaggacaca | 60 |
| ggttacaggg | cctgacgtca | ctaacggtaa | ctgacaatct | tggaatggac | cctactgctg | 120 |
| atgtttcaaa | aggacacaga | ggtgaactgg | tcacttctaa | ttaagaagag | ccagtggggg | 180 |
| gggggaagct | gaaaaccaa | aatccacgta | gacatacgtg | gcagtgtgaa | cgtctgtcct | 240 |
| ccccttcctt | ctcctcactt | cctctcctcc | tcctcactca | ggctggtatt | ctcctggtgt | 300 |
| gcggtatgca | gcttgccctg | cagaagggct | gccagttttt | tagatgtctt | tttgagaaac | 360 |
| gagctgcccg | gatgggcact | gttcacgtgc | aggtacaggt | cctcctgggt | ggggcccgtg | 420 |
| tagccgcaat | cctcgcagac | gtagagcttg | tcctcgccgt | gcttataggc | atactgctgc | 480 |
| tgcaccccat | ggattttctt | cag | | | | 503 |

<210> 1365

<211> 245

<212> DNA

| | | | | | | | |
|------------|------------|------------|------------|------------|-------------|--|-----|
| <400> | 1365 | | | | | | |
| ctgggcggct | ccacgctcat | ccagtgggcc | taggttctga | ctgaccagcg | aacaaaaact | | 60 |
| gtgacagaga | tctaggattt | cattcaggca | gtgaaacacc | tacccgggaa | acagagtttg | | 120 |
| cattaggaaa | ggaaggaagg | tacatccatg | aagttaaagt | gttaggagaa | cagtcctgatt | | 180 |
| aatagctgat | ctaattaata | gctgacctcc | caaatctgac | aggatagaca | ctgccacgtg | | 240 |
| caagc | | | | | | | 245 |

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

```
<400> 1369
ctgaaggcaa tgggggactg aggaaggagg cagcagaagt aggagaggag caagaatcca      60
gaagggaaat gagaacgaca aaactgaagt gcacttcaac atcctgcagc caaaggggta      120
```

```
<210> 1370
<211> 540
<212> DNA
<213> Homo sapien
```

```
<210> 1371
<211> 142
<212> DNA
<213> Homo sapien
```

```
<210> 1372
<211> 377
<212> DNA
<213> Homo sapien
```

```
<210> 1373
<211> 504
<212> DNA
<213> Homo sapien
```

<400> 1373
ccatgctaag tttgggaacc gctggtgatg ggacatggat gcttgcaacc gaccgtgggc 60

```

ggatgtggtt gaccagatgg cagaggacga caccatccat gagggctgcc cccaggtctt 120
cgtgcagact gaccttcaat ctcatctcaa tgctctcacg aagttgttcc accagctctt 180
tctcttctct catctgctcc attttctctc ggattgtaaa ctgcgggtct atagattcca 240
aatttctctg aggtcttaga aacacagact cagaaatcaa atgaggatgt ctcagaaagg 300
agtcactttt ccagaggcag gctgcccctt aactcagccg agcagcagga accactgggg 360
ccaaagctat tttatcttcc ttaggtaaaa aaaaatcaat agaataattt ttccccgctt 420
acatgctccc accactgatg aacgcgatct tcagcaagaa gaactttgag tccctctccg 480
aagccttcag cgtggcctct gcag 504

```

<210> 1374

<211> 201

<212> DNA

<213> Homo sapien

<400> 1374

```

cctccgtaag atgcttgaca attttgactg ttttggagac aaactgtcag atgagtccat 60
cttcagtgtc tttttgtcag ttgtgggcaa gctgcgacgt ggggccaaagc ctgagggcaa 120
ggctataata gatgaatttg agcagaagct tcgggcctgt cataccagag gtttgatgg 180
aatcaaggag cttgagattg g 201

```

<210> 1375

<211> 295

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(295)

<223> n = A,T,C or G

<400> 1375

```

ctgtgaggct gnttccaagg aggaaaacaa ggaaaaaaat cgatatgtaa acatcttgcc 60
ttatgaccac tctagagtcc acctgacacc ggttgaagggt gttccagatt ctgattacat 120
caatgcttca ttcattcaacg gctaccaaga aaagaacaaa ttcattgctg cacaaggacc 180
aaaagaagaa acggtgaatg atttctggcg gatgatctgg gaacaaaaca cagccaccat 240
cgtcatgggtt accaacctga aggagagaaa ggagtgcagg tgcgcccagt actgg 295

```

<210> 1376

<211> 318

<212> DNA

<213> Homo sapien

<400> 1376

```

ccagcgctac tgtactggcc cagggcagag ttcatgtatc tcgtcttgac cacgtctaca 60
ggggaggcga tgacagtggg gcagaagcct gccccaaagg cagaagtga gtaggcaagg 120
aggatcatctg tcatgaggtt ggctttcagg agggcatcct tgatgaggtc ataggtcacc 180
agctcagcac agttgacaat ggcattacga gcaacattgg gggagggtccc tttccagagg 240
ccccggaacc cttcctctcg ggcaatggtc ttgtaggcat tgacggtgct ttggtatctc 300
cgaccacctc cagcccg 318

```

<210> 1377

<211> 143

<212> DNA

<213> Homo sapien

<400> 1377
 gtggattccg ytcggggcac cgatctcgcc aagatcctga gtgacatgcg aagccaatat 60
 gaggtcatgg ccgagcagaa ccggaaggat gctgaagcct ggttcaccag ccggactgaa 120
 gaattgaacc gggaggtcgc tgg 143

<210> 1378
 <211> 98
 <212> DNA
 <213> Homo sapien

<400> 1378
 aaatattggt aataggctcg caacagcaac tatagaagta caactcaata gatggcatta 60
 aaacatattg tagtgtggat atatatTTTT tctTTTTT 98

<210> 1379
 <211> 330
 <212> DNA
 <213> Homo sapien

<400> 1379
 aaagatgttc acgttacgct ggaccaaatt aagacggctt tctccctctt gctgacgtgc 60
 cccagccgtg ataatgacca gcttggagtt tgcagttaca ttatagtctt tgccagagac 120
 aatctttggt gttctaagga aaaggctgcc atgttggaga tccatcatct ctcccttcaa 180
 tttgtcttcg acgacatcaa caagagcaag ttcatctgcc aagtccttca ttaagatact 240
 gatggcacag gccatgccaa cagcaccaac cccaacaact gtaatcttat tctgggggggt 300
 ctgttcttcc tttagaagat tataaatcag 330

<210> 1380
 <211> 269
 <212> DNA
 <213> Homo sapien

<400> 1380
 ccactcctgg aaaccactg atagatgagt ttccccatt cttctggcct ccgccacatg 60
 atcaggaagc tggacttgct cttatccaac cactcgaggt tccctttctt cctcagttcc 120
 tctaatacaa tctggatcga ctccacagga agctttcgtc gtagcttgac gttgttgaag 180
 agcgggctct cctgagcttc catcacgctc atgctggact gtttgtgcag gcggcagaag 240
 gacaggacca gcgagcacca ggcggccag 269

<210> 1381
 <211> 232
 <212> DNA
 <213> Homo sapien

<400> 1381
 aaaagagagg aaaggcagtg cagggctgga ggtcctggag ggtggcggcg ggtcgtccta 60
 actagcaggc tgaaaggctg tggaggggat gccttcactc agaggaagtt cacagccacc 120
 tgccttgga catgtacctg ttcattcttt cgtaatgtta gtattcattt tgctatcttc 180
 ctgttgccat ttccaaacag tgtcagtatg tttttgttaa atacgaacat tt 232

<210> 1382
 <211> 348
 <212> DNA

<213> Homo sapien

<400> 1382

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| aaacgtgcta | aagggaaagg | aatctgacat | tctgggtaaa | tcttactcaa | tctaaatcaa | 60 |
| agcttggttt | tcaggaggag | gaaggtgcga | gcgcaggcag | aggtgctgaa | tactcctctt | 120 |
| ctgattcact | tccatcatcc | tctttctctt | ggctactgcc | ctcagtgcta | agccgggtcaa | 180 |
| acccttttctg | actgtagccc | ttacggcttg | caaagaaatt | accaagggtt | aagcctccac | 240 |
| ttccctttcc | tctaaatctt | cccagtactc | ttcctgaact | cgtctcgagt | ttgtgttcag | 300 |
| aatctccaaa | ggccttgat | tttttccacc | gaataaatat | ggcaatgg | | 348 |

<210> 1383

<211> 293

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(293)

<223> n = A,T,C or G

<400> 1383

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| ctgcttcaan | acctcagctt | catgggactt | gcgtctttct | tctgcagctt | ctaatttctt | 60 |
| ctgaatttcc | tccagggaaa | gatccttctt | ctttggaggg | gaaaggggga | attctggaac | 120 |
| agattctttt | gaccgagggc | tgagaatcag | ctcaaaagcc | tggcccagag | cacgcttctc | 180 |
| cagttctttc | acctggatat | cagaagaagc | catggtgaat | agaagacaag | cgacaggcag | 240 |
| tgtattctgc | acaatcaact | gggataagga | aagtcctgct | cagtcaggagc | cgc | 293 |

<210> 1384

<211> 573

<212> DNA

<213> Homo sapien

<400> 1384

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| ctgaagcaac | ttgggattaa | ttgcttgatt | agcttcacga | agcacagaga | taaggctcgt | 60 |
| cacttgcttt | atgttattag | gtgtaaagaa | agtgtatgct | gtgcctgttt | tggtagctgcg | 120 |
| agcagttctt | ccaattcgat | gaatataatc | ctctgaggag | ttagggtagt | cataattgat | 180 |
| gacaaatttc | acatcttcca | catctagccc | tctggaggcc | acatctgtag | caatcagaat | 240 |
| aggagctttt | ccatgtttga | attcatttag | aaccacgtca | cgtctctgtt | gactcttgct | 300 |
| accatggata | cccattggcag | gccacccatc | tctcctcatt | ttcttggtta | gctcatcaca | 360 |
| tcttcttttg | gtttccacaa | aaacaatggt | tttattctcc | ttctcactca | tgatctcttc | 420 |
| cattagacga | ataagttttt | catccttttc | tacgtcatga | cacacatcca | caatctgaag | 480 |
| aatgtttgtg | tttgactca | gttcaagtgc | accaatgttt | atatgaatat | agtctttcag | 540 |
| gaaatcttca | gcaagctgtc | ttacttcttt | tgg | | | 573 |

<210> 1385

<211> 150

<212> DNA

<213> Homo sapien

<400> 1385

| | | | | | | |
|------------|-------------|------------|-------------|------------|------------|-----|
| ccaaggccgc | tagggctcctt | acccctcagg | atcaactcccc | agccctttcc | tcaggaggta | 60 |
| ccgtctctca | aggtgtgcta | gcagtgggcc | ctgcccaact | tcaggcagaa | cagggaggcc | 120 |
| cagagattac | agatcccctc | ctgtaagtgg | | | | 150 |

<210> 1386
 <211> 159
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(159)
 <223> n = A,T,C or G

| | |
|---|-----|
| <400> 1386 | |
| aaatgatgtt ttggtaaga gtggaccatg agaattagct gacagcatcc cctttctctc | 60 |
| tccctgcctt ggtgggaccc tccctgtgtg accttggtca agtcctcgaa cttttgtccc | 120 |
| gtatttaaga tggagctgnt ttacctactt cataagaca | 159 |

<210> 1387
 <211> 735
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(735)
 <223> n = A,T,C or G

| | |
|--|-----|
| <400> 1387 | |
| ggtgnaattc gcctttgaan ggccgccggg caggtccttt ntgtstgctg aaggcagatc | 60 |
| gcttggtcca caccagctac cactcccagg cagtgcatac ccgccctgtt tgcagaaatg | 120 |
| cacgtgttac tagcatctcc tgggagctga ggcagaccct gtcagttgta tttgatgcct | 180 |
| tcattcacggg gcaggggaaag aaagactggt cctctttccg gatgttctcc cgaacctca | 240 |
| cggagccctg cccctggct tcagagagcc gagtctatgt ggacatcacc acctacaacc | 300 |
| aggacaacga gacattagag gtgcaccac ccccgaccac tacatatcag gacgtcatcc | 360 |
| taggcactcg gaagacctat gccatctatg acttgcttga caccgccatg atcaacaact | 420 |
| ctcgaaacct caacatccag ctcaagtgga agagaccccc agagaatgag gccccccag | 480 |
| tgcctttct gcatgccag cggtagctga gtggctatgg gctgcagaag ggggagctga | 540 |
| gcacactgct gtacaacacc caccataacc gggccttccc ggtgctgctg ctggacaccg | 600 |
| tacctggta tctgcggctg tatgtgcaca cctcaccat cacctccaag ggcaaggaga | 660 |
| acaaaccaag ttacatccac taccagcctg ccagggaccg gctgcaaccc cacctcctgg | 720 |
| agatgctgat tcaga | 735 |

<210> 1388
 <211> 369
 <212> DNA
 <213> Homo sapien

| | |
|---|-----|
| <400> 1388 | |
| ctggggacag cctacagggg cctccagcct gtgccagacg aggaggtgat tgagctgtat | 60 |
| gggggtaccc agcacatccc actataccag atgagtggct tctatggcaa gggtoctcc | 120 |
| attaagcagt tcatggacat cttctcgcta ccggagatgg ctctgctgtc ctgtgtggtg | 180 |
| gactactttc tgggccacag cctggagttt gaccaagcac atctctacaa ggacgtgacg | 240 |
| gacgccatcc gagacgtgca tgtgaagggc ctcatgtacc agtggatcga gcaggacatg | 300 |
| gagaagtaca tcctgagagg ggatgagacg tttgctgtcc tgagccgcct ggtggcccat | 360 |
| gggaaacag | 369 |

<213> Homo sapien

| | | | | | | |
|-----------|----------|-----------|----------|----------|----------|-----|
| aagatg | ctggcatt | ctttttatt | gtaagg | ggtactat | gttattgg | 60 |
| agaaatc | agttttca | tgtatatat | tatagttt | aaaaaga | aaacaacc | 120 |
| gacaaacc | tgatgctc | tgctcggc | tgaggctg | gggaagat | cttttggg | 180 |
| aggctgtag | tcagggcg | cactgtgag | ctggacct | tgactctg | gggggcac | 240 |
| atttagctt | aggttgtc | gtttctgt | atagtga | agcattct | cgccatct | 300 |
| cctgtggc | aagggggg | ag | | | | 322 |

<213> Homo sapien

| | | | | | | |
|-------------|------------|-------------|------------|-------------|------------|-----|
| aaatattagw | tgagacttta | caggcacata | actgttcaga | tagaaacaaa | cataacagac | 60 |
| taaaataactt | tcaaaattaa | agccatctag | aaaatggaag | taactgaaac | tgtagccatt | 120 |
| acaattcttt | ttctggtttt | gagcaaaaat | tttatctctc | tggcaaaaaca | cctttgtctg | 180 |
| atcatttgag | agacagggtt | cttgatact | gtttcttcaa | cgtaaacctc | atttacaaaa | 240 |
| atagtgcac | agcattatga | ataaaactatg | aattggggac | catggaaatg | cactagaaca | 300 |
| aattttgtaa | aaatatggca | gatatggaag | ttaaaaatag | aatggatgca | aggactgtac | 360 |
| taaagggtgt | tggtgtagtt | acaatgttca | ctttgcacaa | ctatccctat | agtctaggta | 420 |
| qccattqggt | ttctctcag | cagtgtcaga | | | | 450 |

<213> Homo sapien

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| aaaaaatcat | aaatgggggt | tcataatcca | aagttgaaac | atttattctt | catagcttca | 60 |
| aaaatttaaca | accaattgta | gaccatgctt | tccaaatcca | gtcttctttg | ctattttttca | 120 |
| aaacttctga | gatctagtat | taaactgctc | cattctaaat | gtatagtttt | agataagtat | 180 |
| tgtacacttg | ttgataaagg | ttttctgaaa | gcagtctatc | aaatataaag | aatgggtttct | 240 |
| atctaagaat | cagcagtgag | ggaagaaata | ttaaacacct | atcaagaaat | caattattca | 300 |
| tttt | | | | | | 304 |

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctggaagaag | aactgagaca | gcagaaagaa | gcagcttggt | tcaaggctcg | tccaaacacc | 60 |
| gtcatctctc | aggagccctt | tgttcccaag | aaagagaaga | aatcagttgc | tgagggcctt | 120 |
| tctggttctc | tagttcagga | | | | | 140 |

<212> DNA

<210> 1397

<400> 1397

<210> 1398

<211> 261

<212> DNA

<213> Homo sapien

<400> 1398

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|-----|
| aaaattataa | ctactcattc | tttcttttagc | cttagataat | ttgagcagaa | gccacaacaa | 60 |
| gcaaaccaca | ataaatntag | aattggcgaga | aatccacatt | aactcctctt | cccaagtttc | 120 |
| cacactacta | ccattttacag | tggtagggtt | gtaatgtata | attatgtaat | gcasaaacta | 180 |
| gctttgactt | gtgttracgat | gcactgtcaa | aggaagcaaa | gtaagaattg | aaattccaca | 240 |
| ttcccgaat | ttaaacactca | g | | | | 261 |

<210> 1399

<211> 195

<212> DNA

<213> Homo sapien

<400> 1399

```
ctgattttat ttccttctca aaaaaagtta tttacagaag gtatatatca acaatctgac      60
aggcagtgaa cttgacatga ttagctggca tgattttttc ttttttttcc cccaaacatt    120
gtttttgtgg ccttgaattt taagacaaat attctacacg gcataattgca caggatggat    180
qqcaaaaaaa agttt                                     195
```

<210> 1400

 $\langle 211 \rangle$ 120

<212> DNA

<213> Homo sapien

<400> 1400

| | | | | | | |
|-------------|------------|------------|-------------|------------|-------------|-----|
| ctgcctccaa | ccctttgggt | ctccaccacc | caagtttctt | gtagggtccg | ccgggtccag | 60 |
| gatacacaggc | ctgggtttcg | tgaagctgct | tctcagggtac | ttttcaataa | tgggggttttt | 120 |

<210> 1401

<211> 284

<212> DNA

<213> Homo sapien

<400> 1401
 ctgtagccaa aaagatgctg gggcagattg tggacaagta gaagcacctc cttccccctct 60
 gcgacattga acggcgtgga ttcaatagtg agcttggcag tgggtgggcgg gttccagaag 120
 gttagaagtg aggctgtgag caggagcctc tgccagggga catgcaatct gcagggaggg 180
 gctgaggggg gtcccatggt ctctgctgtc ttctctgtcc acctctttgt agaggagctt 240
 gagctccagg aatgctcttg tcagggctgc tgtgactggt ggcc 284

<210> 1402
 <211> 198
 <212> DNA
 <213> Homo sapien

<400> 1402
 ccaggtttct gctggtacca ggctaagtag ctggtgctgg cgggaacact gtgactggcc 60
 ctgcaggaga ggggtggctct ttcccccgga gacagagaca gcgtgtctgg agactgtgtc 120
 acttcaagct ctgcgatgcc atctgggagc cagagtagca ggaggaagag aagctgcgct 180
 ggggtttcca tggttccc 198

<210> 1403
 <211> 441
 <212> DNA
 <213> Homo sapien

<400> 1403
 aaactcaaaa ttgacaaatt aactagcttg ctttttgtca tttggaagac taccattatt 60
 caaatttatt atgtaataca ctcatccaga taatgaaaca tctgcgaaaa aaagtgtggg 120
 aatcacctca tctgtgcata aaatggctat tatacatgaa tgcagacgtt tgaagttaga 180
 aaggaatata actcaaatag caaaaggctc taattacaga gtttaccat aagcagtttt 240
 attttcaaaa gtacatagta agtccagact gggctattgc caaagaacta atcttttagtc 300
 tacttcaaca tgttacatgg tattcctgac tctacagact atcagcatct gtggagggtta 360
 gctcctaag gtcccaaaga acaggaaaca tgcaggaata aaggactcct catgaagagc 420
 aggtgggagc gagtgggcag g 441

<210> 1404
 <211> 243
 <212> DNA
 <213> Homo sapien

<400> 1404
 tgaaggggtt cttggaagac ctggcacctc cagagcgcag cagcctaatt caggattggg 60
 aaacatctgg gcttgtttac ctggactata ttagagtcac tgaaatgtc cgccatatac 120
 agcaggtgga ttgctcaggt aatgacctgg agcagttaca catcaaagtg acttcaactgt 180
 gcagtcggat agagcagatt cagtgttaca gtgctaaaga tcgcctggct cagtcagaca 240
 tgg 243

<210> 1405
 <211> 168
 <212> DNA
 <213> Homo sapien

<400> 1405
 aaaccactgg atctatctaa atgccgattt gagttcgcga cactatgtac tgcgtttttc 60
 attcttgtat ttgactatct aatcctttct acttgtcgtc aaatataatt gtttttagtct 120
 tatggcatga tgatagcata tgtgttcagg tttatagctg ttgtgttt 168

<400> 1409
ccaqtccaac ctgctcctca ttattgtata aatgagcaga atcaatatgg cggaagccag 60

```

cttcaattgc caatttgggtg gcctctaaag ctttactttt aggaacctct gcaggcgcac 120
aggtgccaaa tcccaggaca ggcattgaagt gaccatcatt cagcttcaca cactgatatt 180
tcgaatccat ttctgtcact agcctggc 208

```

```

<210> 1410
<211> 404
<212> DNA
<213> Homo sapien

```

```

<400> 1410
aaaaaaagga aaaagtttta ttacgaaact agtttgtata aaacagggtt atacatattt 60
ttgtaagttt gtaataaaac agtaagaaaa aaaaggcagt aatagaaatc tccaaaaggc 120
aacctatcaa aaccaactgg ctgccacttt gagtttggac agtagctgca taaactttgt 180
tcttcttgar cagtatttaa taacatcatt aatacattaa caacatttct ataaagtaag 240
acacattggt gctgaagtac aactgggtgg ctcttgatct cacctatgag gagagttctt 300
tacamawcca catagggaaa attgcagttg taagggtgarc tacacatcta aaatatgcag 360
aggtaatagc attacatggt aaagtatcaa gatatacaca tttt 404

```

```

<210> 1411
<211> 623
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(623)
<223> n = A,T,C or G

```

```

<400> 1411
ccacttggtg agatatgggg agcctacact ccggaggggst gtaccttttag cactggccct 60
catctctggt tcaaatccac gactcaacat cctggatacc ctaagcaaatt tctctcatga 120
tgctgatcca gaagtttcct ataactccat ttttgccatg ggcattggtg gcagtggtag 180
caataatgcc cgtctggctg caatgctgcy ccagtttagct caatatcatg ccaaggaccc 240
aaacaacctc ttcattggtg gcttggcaca gggcctgaca catttaggga agggcaccct 300
taccctctgc ccctaccaca gcgaccggca gcttatgagc cagggtggccg tggctggact 360
gctcactgtg cttgtctctt tcttggtatg tcgaaacatt attctaggca aatcacacta 420
tgtattgnat gggctggtgg ctgccatgca gcccgaatg ctggttacng tttgatgagg 480
agctgcggcc attgccagt tctgtccgtg tgggccaggc agtggatgtg gtgggcccagg 540
ctggcaagcc cgaaaactat cacagggttc cagacgcata caaccccgat gttggtgggc 600
ccacggggaa cgggcagaat tgg 623

```

```

<210> 1412
<211> 171
<212> DNA
<213> Homo sapien

```

```

<400> 1412
gcggcgctgg ggggtgctgga gtccgacctg ccaagtgccg tgacacttct gaaaaatctc 60
caggagcaag tgatggctgt aactgcacaa gtgaaatcac tgacacaaaa agttcaagct 120
ggtgcctatc ctacagaaaa gggctctcagc ttcttgggaag tgaaagacca g 171

```

```

<210> 1413
<211> 189
<212> DNA

```

| | | | | | | |
|-------------|-------------|------------|------------|------------|-------------|-----|
| cttgattttag | gatctgtggg | gcagggcaat | gtttcaaagt | ttagtcacag | cttaaaaaaca | 60 |
| ttcagtggtga | ctttaatat | ataaaatgat | ttcccatgcc | ataattyttc | tgtctatttaa | 120 |
| atgggacaag | tgtaaagcat | gcaaaagtta | gagatctgtt | atataacatt | tgttttgtga | 180 |
| tttgaactcc | taggaaaaat | atgatttcat | aaatgtaaaa | tgcacagaaa | tgcatgcaat | 240 |
| acttataaga | cttaaaaaat | gtgtttacag | atggtttatt | tgtgcatatt | tttactactg | 300 |
| cttttccctaa | atgcatactg | tatataattc | tgtgtatttg | ataaatattt | cttccctacat | 360 |
| tatatTTTTTA | qaatatTTTCA | qaaatataca | tttatgtctt | tatattgtaa | taaatatgta | 420 |

catatctagg tatatgcttt ctctctgctg tgaaattatt tttagaatta taaattcaca 480
 tgtcttgca gatttcacat gtataccttc aaattctctg aaagtaaaaa taaaagtttt 540

<210> 1417

<211> 350

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(350)

<223> n = A,T,C or G

<400> 1417

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ttnatcatct | aactgtggga | tctatttcat | ttctggaaat | aacacaactt | agttctaggg | 60 |
| ctttcatgca | catgaaatat | aaaacagctt | agttgttctg | aaaacatgac | aatggttaat | 120 |
| tttattcaag | tccaacact | gagttcagag | cacttctcca | taggccccat | taatctctcc | 180 |
| aggtttctgg | gagtatcatt | aaatccctcg | gcattcctta | gaagcaggtg | cttagcaaac | 240 |
| atccagtttc | caaagtagag | tcagaggggc | ttgatcctga | aagtgtagta | ttttcctgcc | 300 |
| ttgtctact | ggtatagctt | cttggaccta | aaatctctct | cctgctgagg | | 350 |

<210> 1418

<211> 425

<212> DNA

<213> Homo sapien

<400> 1418

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| tgctaggcag | ccttattttc | ataacccawt | tagggaaagg | aaatttagga | ttttcaaggc | 60 |
| tacattaatt | tttctccat | caaactctga | tttgttcttg | ataaaaatga | gttcttttgg | 120 |
| ggaaattctt | tctttagaca | ccaacttggg | ttttctcatc | ttccacagaa | taattgaacc | 180 |
| cctgacctct | agatgttcaa | aattccgctt | caagcctctg | tcagataaaa | ttcaacagca | 240 |
| gcgattacta | gacattgcca | agaaggaaaa | tgtcaaaaatt | agtgatgagg | gaatagctta | 300 |
| tcttggttaa | gtgtcagaag | gagacttaag | aaaagccatt | acattttctt | aaagcgctac | 360 |
| tcgattaaca | ggtggaaagg | agatcacaga | gaaagtgatt | acagacattg | ccggggtaat | 420 |
| accag | | | | | | 425 |

<210> 1419

<211> 390

<212> DNA

<213> Homo sapien

<400> 1419

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaactcttgc | tattgaattg | agatgattaa | aatggtgact | taatccgtag | ttattttgca | 60 |
| cccactgaaa | ggaaagtgc | ttccagaata | atatgaagta | tctaaaagtg | tcaccttttc | 120 |
| ttgcctgac | aacaatttgg | gcttcctgtt | tgtacaaggg | gccatttggc | atacctttca | 180 |
| cagcttttat | caggccaagt | taaaggctga | ctacattttt | tcacatgag | gaaagcagtt | 240 |
| gaaatgaggc | atgagttact | gtgcattggg | attttagaac | aattttcttg | tgacagctct | 300 |
| tttgtgaag | ttaggttctt | aaaagtgcc | atgatggtca | cttaaaatgt | gcagtaatag | 360 |
| cactgccagg | atcaagcatg | aaaggctttt | | | | 390 |

<210> 1420

<211> 480

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(480)
 <223> n = A,T,C or G

<400> 1420
 ttgctgaaca atgacatcgt tttctccagg ggttgaaatc catgtccatg gctgacaacc 60
 caacaaggct gggacccaaa ttctgtacaga gatgaggcag agtggagaga aacaactctg 120
 gctgagccag agtctccagc cactacttct tattcctggg ctttagctct tcggctgcat 180
 tacgcaggaa aatgtaattt tttttctggg gattataaaa ttcattgtccc tttgaccagt 240
 cgtagctgga agcgtatgca aatatgtttc cattgygatt gaaacagcaa gctgasatgg 300
 gctgayctaa ctgttccgaa gnttttagtt ttgktctggc atctttgycc cagaagctga 360
 atctaccatc agatcccaca gttgcaaggg tgccatgaac aggatggaac gccgattcca 420
 tttaccgcga taaatgycct gaggagctga agtggttggtt ccattagatc gatgacattt 480

<210> 1421
 <211> 453
 <212> DNA
 <213> Homo sapien

<400> 1421
 aaactgattg aggtcacagt attttattat ttggggtoct caccacagga aacactgcga 60
 tacaggggca aaagagatgg cagtgccaat taaattaata caacaaaatc aatgcagcac 120
 caaccaagac tgccaggctc ggtgtcatgg gtatgccag agcccaggag ttcagaaggg 180
 ccctaagcct gatttaatgc tctgctgttg atgtcttgaa attcttaaca atttttgaac 240
 aaggggcctg cgttttcact tcgcactggg ccttgcaaat tacatagcga gtgctcataa 300
 aagaactcag aaacgtggta cctctcttcc tgggtggatac aaataaagaa atctggatcc 360
 aaagttgaaa gttgctggcg atatcattca agtaggactc taaatagtgg attaagatga 420
 ggggtgggcct ggggtgaagat tctttccagc ttt 453

<210> 1422
 <211> 542
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(542)
 <223> n = A,T,C or G

<400> 1422
 ttttcttgac cactatacgg cacaacctag gggstgtawa aaacctascr caatgcagaa 60
 ggggtgaagct tcatgacaat tggctcgcgc aataatttgg gggatgtaac atcaacgaat 120
 cagacaacaa aagcaaggga atacacatgg nactaaatca gtgtgnggaa aaatatccca 180
 aacaggcaaa gcacaacatg gamtagatat atgcacattn atggaccctg naggcakkac 240
 tcacaaacat actacctggg aagcamctgg acctttaagg gatgaggtag attcaacaaa 300
 cagggcancg tatmttcac tgggtagca ttccagcctt aaaaataang aaatcttgaa 360
 aagnactaca ataaggacaa atctcgaaca cattctgtta agtaaaacaa gacaagccaa 420
 aaagggaana ctgtataatt acacctatgt aaaatattta gtcaaaactca aagaaaccaa 480
 gtgttgtagt ctcagcaggg caccaagatg naaacagtct ctcatagnct gagatangca 540
 tc 542

<210> 1423

| <400> 1426 | | | | | | | |
|-------------|-------------|-------------|-------------|--------------|-------------|--|-----|
| gtggtagaaa | gagatgacgg | aagcacatta | atgggaaatag | atggcgataa | aggcaaacaa | | 60 |
| ggcgggtccca | cctactacat | agatactaata | gctctgcgtg | ttccgagggg | gaatatggag | | 120 |
| gccattttcac | ctctaaaaaa | tgggatgggt | gaagactggg | atagttttcca | agctattttg | | 180 |
| gatacatacct | acaaaatgca | tgtcaaataca | gaagccagtc | tccatcctgt | tctcatgtca | | 240 |
| gaggcacccgt | ggaataactag | agcaaaagaga | gagaaactga | cacaggttaata | gtttgaaacac | | 300 |
| tacaacatacc | ctgccttctt | cctttgcaaa | actgcagttt | tgacagcatt | tgctaattggt | | 360 |
| cqttctactq | qqctqatttt | qqacagtgga | gccactcata | ccactgcaat | tccagtcacc | | 420 |

```
<210> 1427
<211> 144
<212> DNA
<213> Homo sapien
```

```
<210> 1428
<211> 214
<212> DNA
<213> Homo sapien
```

```
<210> 1429
<211> 253
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|-----------|------------|------------|------------|------------|------------|--|-----|
| <400> | 1429 | | | | | | |
| ccactagtc | antttngtg | aattctgaag | ccttaattgc | ttatatccat | gtttctagtg | | 60 |
| aaatgagag | gtataacaaa | aaagagaaca | ggaggaaagc | ttcgctgtgc | ctgaggaaat | | 120 |
| aatctagtc | aggcagcaag | tctggatagt | gctatagaga | tgagatacct | gagcagttcc | | 180 |
| agaggaaag | gtggagatca | gaggccagtt | ttcagtgaa | actgtaaaga | aaagccagat | | 240 |
| gatgttcct | gga | | | | | | 253 |

| <400> 1430 | | | | | | |
|-------------|------------|------------|-------------|-------------|------------|-----|
| aaatttttact | agtgttactt | aatgtatatt | ctaaaaagag | aatgcagtaa | ctaatgccct | 60 |
| aaatgtttga | tctctgtttg | tcattacttt | ttcaaaaatta | ttttttttctg | taaagtataa | 120 |
| tatataaaaac | ttcttgctta | aattgaattt | ctatatattgt | ggttaaatgc | agtttattaa | 180 |
| agggatcatt | atcagtaatt | tcatagcaac | tgtttcagtg | tttttggttt | tt | 232 |

<210> 1431

<211> 734
 <212> DNA
 <213> Homo sapien

<400> 1431
 cattatacaaa cactatatattg ccagggtcaaa gagggcaggg acgtaaatgt acactaaaat 60
 gcmaatgtat cccaaagaga taaaacaaat tccattttaca gcatgaaggt ttacaaatgt 120
 acacctgtac aaccaaggaa agcatcacta ctaaattagc aaggctttta taataaacat 180
 tgaaasaaga tttcctttca aagtgtaaac ttacatctat tactacacac acaatgcata 240
 tattttataga aagcaaaaag agctatctga atatgtaatc atgcttaaat gctgagctat 300
 caaattcact tttcagtggc cccttttcat ctctatctgg ttctactttt ctgcctctat 360
 gaaaaagcaa aataaagctc aacacttctt caacatgtct gtaattctat aagcaaaaca 420
 aaatacaaat ttccactctt tctcattgca aaccaaactg aaaagttaat aagtgactta 480
 acttttctatt tagtgacttt aattggaagt gtcaccatga ttttgtattt aactcttaca 540
 acaattacat atgtaagtat atacaatatt tctgtacatt gccagagaca ttttagggca 600
 gtaattgtat taaaaccaca tctactgtaa ataatgttag gttctttttca tctcaaacca 660
 ctttattctt gccacttac tcgttatattg catgatagtt tgtgaattat caaaatacaa 720
 cttaactctt taaa 734

<210> 1432
 <211> 542
 <212> DNA
 <213> Homo sapien

<400> 1432
 tttaagaaag agccttttgag aaacatgcat actttttctct tttctctat attcaatact 60
 catatagcct aaaagatgga aactgggttca agaatttaaa tgacttggtc cctaaaaagt 120
 taatctcttc accctttgtga aatatatcaa gtgcttttcta taaataaggg caggaaatgc 180
 taacttcata agcatagtcc tagtcattaa aataatttga tcatcttcta aattttaagt 240
 atgatatgaa cacagtaata tggaaaatct caatatactt aacacttctt aaacagcaca 300
 atgaaatggt gttcaaggtc tgaatttaatt tgctacagga cctaagcaag tctgtttgct 360
 tatcttttgg ctttaaaatt ctttaagtct aaaatgggtga taattttaga ataaactgac 420
 aatgtgggga acaaacttaa attcacaac actaccata tgetcaaaaa ctctctggga 480
 taattagttt cttcattgta actattgatg tactattatt tcatctttcc attagctcta 540
 ct 542

<210> 1433
 <211> 175
 <212> DNA
 <213> Homo sapien

<400> 1433
 tttaattgat tcaaaaaaac ttgacacctg tcatgtaggc caaaaaatag tagcgaacta 60
 tactaagtgg tatagcccac tgtggagtgt ggtcttttac tcttccaaat agcccaagtt 120
 ggcaaagggt acttaaaaaac ctgccccca aaaagctaac ttttggtaga ttttt 175

<210> 1434
 <211> 90
 <212> DNA
 <213> Homo sapien

<400> 1434
 ttaatcacta ttgatggaag cttatatctc ttatgaatat atacatgtat gcatatatac 60
 atctctgtat gaatcactca aagcaatttt 90

<210> 1435
 <211> 153
 <212> DNA
 <213> Homo sapien

```
<400> 1435
tttacctttg tgctttgaag gttctaccat ttakaaagta aaaagccaac ccacagaatg      60
gaagaaaaga ggacagactc taacaagcgt tcacaaagat ggagagaaat tgtaaccctc     120
atatattgct ggtagaattg tagaaagatg cag                                     153
```

<210> 1436
 <211> 483
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(483)
 <223> n = A,T,C or G

```
<400> 1436
tttttagttt aaagaagagt tttgccactt aracanggga gctwtgtctg gaaaatacac      60
tgagttgaaa cacttcaccc ttggaaggat tatataagat gaacagytgt gataaatgtg     120
tagattagag ggatgtgaat gggcagttag tccagtgcc ccatTTaaga ggccaagatc     180
ctgattcaga ggaggcatcc tttgcccaga gctgcttagc taatctgacc aaatgTtggg     240
aaaaatgtct cacctaaccc actattcctt aattatggat tttgtgaaaa acaatagaac     300
atgttaatga gtaatttata ttagttcgat gtattacaat tttttagctt taaattacag     360
ytttcttata atgttgaaat gttttagaat cctttgaatc taagtatttg tttcctaaat     420
gaaacatttg tacaacattt gatgttttta cttatgaaat attctcctcc cccaagaaaa     480
ttt                                                                483
```

<210> 1437
 <211> 171
 <212> DNA
 <213> Homo sapien

```
<400> 1437
ttttgccacc tcaagaagcc attttcttgt ctgtttcctt ctttacctac ccctacaacc      60
tatgaacaaa taccataact taaaaattta ggtagtctac aactcctaca aatttttaagt     120
tcagagacta cccaagaac tgtggaagat gcagcaatat aaaagttttt t                                     171
```

<210> 1438
 <211> 408
 <212> DNA
 <213> Homo sapien

```
<400> 1438
tctgagtgga ggtaggctaa caacacattt tgactttstc ctcaaaggat agctttgaaa      60
aacaagtgta accaattgtt acaccaaat aaaatggcaa tattaatcg gtaacaaaac     120
gatccacatt ttatacaata ttgtatttcc aaacatacat aggtcatgaa aatcagagaa     180
cctaatatag caccgttgaa accattcatt atccttcatt tgtgtatgca attcagaatt     240
tcggcagaag acaacaaatg gaaaatgcct ttcgtttcta taaatcattt tggatttcaa     300
ttaaatcttt gccttagtaa agggatttct tatctcaaga tcaattagcc gtttttagct     360
```

ccaccgtttt ggaagtaaaa atgatgagct acatctactt ttttaattt

408

<210> 1439

<211> 168

<212> DNA

<213> Homo sapien

<400> 1439

| | | | | | | |
|-------------|------------|------------|-------------|------------|------------|-----|
| ttacacaaca | gctataaacc | tgaacacata | tgctatcatc | atgccataag | actaaaacaa | 60 |
| ttatatattag | cgacaagtag | aaaggattaa | atagtc aaat | acaagaatga | aaaacgcagt | 120 |
| acatagtgtc | gcgaactcaa | atcggcattt | agatagatcc | agtgggtt | | 168 |

<210> 1440

<211> 307

<212> DNA

<213> Homo sapien

<400> 1440

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| tttcacatac | gaagaaatca | actgtgatta | tgaagtgaca | gccagctaaa | tatgtcttgt | 60 |
| atcttctctc | ttcctttttt | tgctaaactc | atcctttact | tccattcctg | cttccatggg | 120 |
| aatgcaggct | caaataaatt | actaggatac | aagattactt | caagcctctt | ttctgtggaa | 180 |
| ctcataatat | gataagcatt | tgttacaaga | ttgcctgtag | ttgtttaggg | gacaaattat | 240 |
| attagggaaa | gaaagtcttt | cttttagttg | ttaaattttc | tattataatt | gggtactaaa | 300 |
| tttattt | | | | | | 307 |

<210> 1441

<211> 684

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(684)

<223> n = A,T,C or G

<400> 1441

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|-----|
| ttaagttctg | gagtgttcac | ttctgagcct | gaattccctc | ccctgcaaaa | tgggggaata | 60 |
| ccctcctcag | aggggtccctg | cgagggtgag | gggagattca | gcatggcagg | tgtgctgggc | 120 |
| acggcagggc | ctgggaagg | cagatccttt | ccccatccct | gccacaaaca | acccaaacct | 180 |
| ttaaaggaga | gcaatggcct | tgtgtcaaaa | acaaaaacaa | aacaaaaccc | tgtcctagga | 240 |
| gactggggcc | ctaatttcta | atagcaagcc | tttatgagtc | cctaacactc | tactgggctg | 300 |
| agtatctcac | acgccagagg | ataacctgcc | ttctgctcac | caccaccccg | tagtagttgt | 360 |
| cattgtgtcc | atttcacaga | tgaggcaaag | gctcagaaga | gtcatgtgtt | aaaccagctt | 420 |
| ctagagccca | tgcaggagct | gcagggtggga | gaatcacctc | taggtgctct | tcccatagaa | 480 |
| tcctcacctc | ctgagtgtca | ctcactcagc | ttccaatggg | tgtgtgacct | ttgaccagct | 540 |
| ttcttctctc | ctgggcctca | gtttccacc | tggacaaagt | aagaggtctc | ttggcttcan | 600 |
| gtaagttctt | cctaaacttc | tttttccttt | tcatttgagc | atcctcttca | tttttgccac | 660 |
| ctctctgtca | tttacaggct | tttt | | | | 684 |

<210> 1442

<211> 166

<212> DNA

<213> Homo sapien

<400> 1442
 aaaaaatcag cccctaattt ctccatgttt acacttcaat ctgcaggctt cttaaagtga 60
 cagtatccct taacctgcca ccagtgtcca cccctccggcc cccgtcttgt aaaaagggga 120
 ggagaattag ccaaacactg taagctttta agaagaacaa agtttt 166

<210> 1443
 <211> 194
 <212> DNA
 <213> Homo sapien

<400> 1443
 tttgccctgt caaaagaaga gctaaagaca gttatataaa aattaagggtg ggctttcaga 60
 ctggctaaca caacaacatt ccatgagtag atggtaattt atttttgttt atccatttcg 120
 ttgggagcaa ggacaaaaat gtaaattctac accttgctta tcaaaattgc cgaaaaaaga 180
 atgctctgcc tttt 194

<210> 1444
 <211> 96
 <212> DNA
 <213> Homo sapien

<400> 1444
 gagagtcgag agtgggagaa gagcggagcg tgtgagcagt actggggcct cctctcctct 60
 cctaacctcg ctctcgcggc ctacctttac ccgccc 96

<210> 1445
 <211> 365
 <212> DNA
 <213> Homo sapien

<400> 1445
 gggatgagct gaccaagaac caggtcagcc tgacctgcct ggtcaaaggc ttctatccca 60
 gcgacatcgc cgtggagtgg gagagcaatg ggcagccgga gaacaactac aagaccacgc 120
 ctcccgctgt ggactccgac ggctccttct tctctacag caagctcacc gtggacagga 180
 gcaggtggca gcaggggaac gtcttctcat gctccgtgat gcatgagggg ctgcacaacc 240
 actacacgca gaagagcctc tccctgtctc cgggtaaatg agtgcgacgg ccggcaagcc 300
 cccgtcctcc gggctctcgc ggtcgcacga ggatgcttgg cacgtacccc gtgtacatac 360
 ttccc 365

<210> 1446
 <211> 386
 <212> DNA
 <213> Homo sapien

<400> 1446
 tctggaaagt tcttgctcgg gtcccttcac ctccccgccc tttcttarag tgcagttctt 60
 agccctctag aaacgagttg gtgtctttcg tctcagtagc ccccaaccca ataagctgta 120
 gacattgggtt tacagtgaac ctatgctatt ctacagccctt tgaaactctg cttctcctcc 180
 agggcccgat tcccaaacc ccatggcttcc ctcacactgt cttttctacc attttcatta 240
 tagaatgctt ccaatctttt gtgaattttt tattataaaa aatctatttg tatctatcct 300
 aaccagttcg gggatatatt aagatatttt tgtacataag agagaaagag agagaaaaat 360
 ttatagaagt tttgtacaaa tggttt 386

<210> 1447

<211> 261
 <212> DNA
 <213> Homo sapien

<400> 1447
 aaaattataa ctactcattc tttcttttagc cttagttaat ttgagcagaa gccacaacaa 60
 gcaaaccaca ataaatttag aattggcaga aatccacatt aactcctctt cccaagtttc 120
 cactacta ccatttacag ttgtaggttt gtaatgtata attatgtaat gcagaaacta 180
 gctttgactt gtgtaacgat gcactgtcaa agtaagcaaa gtaagaattg aaattccaca 240
 ttcccagaat ttaacactca g 261

<210> 1448
 <211> 404
 <212> DNA
 <213> Homo sapien

<400> 1448
 aaaaaaagga aaaagtttta ttacgaaact agtttgtata aaacaggggtt atacatattt 60
 ttgtaagttt gtaataaaac agtaagaaaa aaaaggcagt aatagaaatc tccaaaaggc 120
 aacctatcaa aaccaactgg ctgccacttt gagtttggac agtagctgca taaactttgt 180
 tcttcttgaa cagtatttaa taacatcatt aatacattaa caacatttct ataaagtaag 240
 acacattggg gctgaagtac aactgggtggc ctcttgatct cacctatgag gagagttctt 300
 taaaaacca catagggaaa attgcagttg taagggtgaac tacacatcta aaatatgcag 360
 aggtaatagc attacatgtt aaagtatcaa gatatacaca tttt 404

<210> 1449
 <211> 230
 <212> DNA
 <213> Homo sapien

<400> 1449
 aaaagttcta gtggtacggg aggagctttg caggaagttt gcaaaagtct ttaccaataa 60
 tatttagagc tagtctccaa gcgacgaaaa aaatgtttta atatttgcaa gcaacttttg 120
 tacagtattt atcgagataa acatggcaat caaaatgtcc attgtttata agctgagaat 180
 ttgccaatat ttttcaagga gargcttctt gctgaatttt gattctgcag 230

<210> 1450
 <211> 194
 <212> DNA
 <213> Homo sapien

<400> 1450
 aaaaactcct tttggtttac ctggggatcc aattgatgta tatgtttata tactgggttc 60
 ttgttttata tacctggctt ttactttatt aatatgagtt actgaagggtg atggaggtat 120
 ttgaaaattt tacttcata ggacatactg catgtaagcc aagtcatgga gaatctgctg 180
 catagctcta tttt 194

<210> 1451
 <211> 106
 <212> DNA
 <213> Homo sapien

<400> 1451
 aaagatgaca aatactggtt aattagcaat ttaagaccag agccaaatta tcccaagagc 60

atacattctt ttggttttcc taactttgtg aaaaaattg atgcag

106

<210> 1452

<211> 349

<212> DNA

<213> Homo sapien

<400> 1452

| | | | | | | |
|-------------|------------|------------|-------------|-------------|-------------|-----|
| ctgcagatcc | tgcggaacgt | caccaccac | gtttccgtga | ccaagcagct | cccaacctca | 60 |
| gaagccgtgg | tgtctgctgt | gagcgaggcg | ggggcgctctg | gaataacaga | ggcgcaagca | 120 |
| cgtgccatcg | tgaacagcgc | cttgaagctg | tattoccaaag | ataagaccgg | gatggaggac | 180 |
| tttgctctgg | aatctgggtg | tggcagcatc | ttgagtactc | gctgttctga | aacttacgaa | 240 |
| accaaaacgg | cgctgatgag | tctgtttggg | atcccgctgt | ggtaacttctc | gcagtcctccg | 300 |
| cgcgtgggtca | tccagcctga | catttacccc | ggtaactgct | gggcattta | | 349 |

<210> 1453

<211> 302

<212> DNA

<213> Homo sapien

<400> 1453

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaaaataatg | tgcaagagca | tcatgagaaa | gaagaggggt | gaagagataa | tccagaggaa | 60 |
| catcaaagt | aagagtatac | actcaaagac | aggtttaaga | aagaccagtc | agagaagtaa | 120 |
| agaaaaaat | caagcaagaa | taatgttgca | aaaattaaca | agaaagttgc | aagcccagag | 180 |
| tggttagcaa | tgccaaacta | ccatgagtaa | gccacataaa | acaagaactt | tgggttcaac | 240 |
| tgctttaaca | atcagacctt | tagattcaca | taacaggagt | tacaaaatta | agagcctctt | 300 |
| tt | | | | | | 302 |

<210> 1454

<211> 268

<212> DNA

<213> Homo sapien

<400> 1454

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| caagcgtaaa | ccgcgggagc | cgagcccagc | taggaatgca | gacctcctga | aaaccaagcc | 60 |
| gaggactgcg | gggtccggtg | tccacgcaga | gtgtcagctt | cctctggtgc | aaccagcaag | 120 |
| tcttcagta | tgaatccac | agaaaccaag | gctgtaaaaa | cagaacctga | gaagaagtca | 180 |
| cagtcaacca | agccaaaaag | cctacccaag | caggcatcag | atacaggaag | taacgatgct | 240 |
| cacaataaaa | aagcagtttc | cagatcag | | | | 268 |

<210> 1455

<211> 207

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(207)

<223> n = A,T,C or G

<400> 1455

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtcgagag | cagccctgcc | caagawtgnc | gggtgggggc | tggtgccaac | gggttcccaa | 60 |
| ggscctttcm | actttkgaak | ggctggartt | cttgggaaac | cmaaacsctg | actacctgsc | 120 |
| ttttttcttg | ggcatygacs | tgcttcattt | ccaaaratga | tggkgcaggt | gaccttttcc | 180 |

207

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<223> n = A, T, C or G

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| aaagaatgca | taccagaaca | tttataagca | gtggagtgg | kthtattaag | aatagtacta | 60 |
| ctacaataaa | cgctggctaa | ataagaagt | cattatgtga | agcactatgg | gtgggtatatg | 120 |
| cttwgmcaca | tactctkgtt | accttgaggy | agatmacrca | tgkgaaccaa | cttcggcata | 180 |
| cattttcagt | tgctgcgagg | aatcatgtgt | tttaacgaaa | tgcgctcagta | tgaaaaactt | 240 |
| gaaaatatct | atgaatgawg | aacgcmttag | gaaaaaaata | kstattctca | tgcaattatg | 300 |
| tacagtctca | ctgtgtarat | ctcaaggcaa | ggtttgcttc | ctgtaaacca | gatcaagggtg | 360 |
| ctatgagaga | ncgcctgnc | ttattgcatt | tcttttctcc | tmctgcgcga | gcatttatatt | 420 |
| qctctagmct | ttatttttgt | gtgcacactg | acatgccatt | aaaratgang | ractatctca | 480 |

```
catgtagaaa argaaagnmc ttgganketa cctcaggtcg ctaccacgct aaggggyaat 540
tctgcaggat atccatcaca ctggcgggcgc gattg 575
```

```
<210> 1460
<211> 444
<212> DNA
<213> Homo sapien
```

```
<400> 1460
ctggggggttc cttccttcac gttgagaacc tggagcagag agtctacca cttagaagaat 60
attagaaaga gttcagcaaa cagagtgagc tgaagtctaa tcctagaagt aaatccattc 120
ctacaagtca tcagcatcac ttgggagcct gttagaaagg caaattcctg gttcagccta 180
acacctacta aatcagaaac tctgggggcg gagcgagca atctgtactt tcacaagccc 240
tgcaggtgat tctgagcctg taaaatttga gaaccagagc tgtccccag gagataaatt 300
aacttctact tttttttgag ctactgcatt ttgggatcct attgttttat cagcttaaca 360
tgcacacctga tatgattact caggatggt tcaaccaatg ttgggtaatg tattatcccc 420
aggaacttat tactagagga gcag 444
```

```
<210> 1461
<211> 536
<212> DNA
<213> Homo sapien
```

```
<400> 1461
ctgcaaccct gggactgacc gggaggctct gattatttac ccmaccacag gtaggttgtg 60
ttctgaatct caggttcaca ggtaagggt cagcatcctc atcctccacg gggttggagt 120
tggtgctggt gatgaagggt ttgggtggct ctgcatagac tgtgatcgtc gtgactgtgg 180
tcctattgag gccactggct gagttattgg cctggcaggt atagagtccg ctgttcttct 240
cagtgatgtt ggagataaag agctcttgtg tgtgttgcct gatgttcca tcaatcagcc 300
aagaatactg tgcaggtggg ttagaggctg catggcagga gaggctgagg ttcaccctg 360
gacggtaata ggtgtatgag ggggaaatgg tggggkrtc ygggccatag aggacattca 420
ggatgactgr gtcgctgtgs tyarcactta atkcgttctg gattccacac tcatagggtc 480
ctacatcatt ccttgtgaca ytgartagag tgagggtcct gttgtcattg gacagm 536
```

```
<210> 1462
<211> 409
<212> DNA
<213> Homo sapien
```

```
<400> 1462
ctgakagacc aggagaagtt ccagatgcag agactgtgat gctcttgact atggaattat 60
tgcggccagt agccaagtta gagacaaaac aggcataagg cccgttatta tttggcgtga 120
ttttggcgat aaagagaact tgtgtgtgtt gctgcggtat ccattgata cgccaagaat 180
actgcgggga tgggttagag gccgagtggc aggagaggtt gaggttcgct cccgaaaggt 240
aagacgagtc tgggggggaa atgatggggg tgtccggccc atagaggaca tccaggggtga 300
ctgggtcact gcggtttgca ctactgagt tctggattcc acatacatag gctcttgctg 360
catttcttgt gacattgaat agagtgaggg tcctgttgcc attggacag 409
```

```
<210> 1463
<211> 502
<212> DNA
<213> Homo sapien
```

```
<400> 1463
```

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccttcagcct | ggatccttta | tattaagatc | aatgaggacc | atttctggaa | gatgtctggc | 60 |
| atggtacaga | ctgtctgagg | ccractgaac | acaggccctt | accctgattt | tatcagtga | 120 |
| aagctatggg | actagtttcc | ttacctctaa | aatggagaga | ataatagaat | cttccgtcta | 180 |
| agactkctgt | gagcataagc | cgagaaaatg | gaggtaaact | gcttagccca | atacttggat | 240 |
| tatcgtaa | attcagtaaa | actagccacc | gttggtattg | taattattat | tttgtatttt | 300 |
| attatacatt | tcatggaaac | ttaaaagtta | gtgataatca | cctcattttc | agttgccttg | 360 |
| ctttcttct | gtaaatttta | ttctctctta | tcttgetcac | tgtctttaag | cattgccagt | 420 |
| ttagtataat | tattttcccc | tatcctctat | aaaatcatat | acaggatgga | tttgttgatc | 480 |
| tcagacatgt | tcactgagtt | tt | | | | 502 |

<210> 1464

<211> 294

<212> DNA

<213> Homo sapien

<400> 1464

| | | | | | | |
|-------------|------------|-------------|------------|-------------|------------|-----|
| ggcggctcgg | actgagcagg | acttttcctta | tcccagttga | ttgtgcagaa | tacactgcct | 60 |
| gtcgccttgct | ttctattcac | catggcttct | tctgatatcc | aggtgaaaga | actggagaag | 120 |
| cgtgcctcag | gccaggcttt | tgagctgatt | ctcagccctc | ggcctaaaaga | atctgttcca | 180 |
| gaattcccc | tttccctcc | aaagaagaag | gatctttccc | tggaggaaat | tcagaagaaa | 240 |
| ttagaagctg | cagaagaaag | acgcaagtcc | catgaagctg | aggtcttgaa | gcag | 294 |

<210> 1465

<211> 249

<212> DNA

<213> Homo sapien

<400> 1465

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| gtgcaggtct | tcagccgtga | cccgttacc | cagctctaag | ggaggtggca | gcatcaaagg | 60 |
| ctccccctgc | ctgcgtggca | gcagggaat | cttgctgcta | cggggcctag | agtcattggga | 120 |
| tctgggggag | ccaccctcgg | gggcaagtgt | ctgccctggg | gctgtacctg | ccttgttttc | 180 |
| acagcgggtga | cccgaagaga | cagcctgagg | tccgtcctca | ctcactgtgt | ttgaggaact | 240 |
| gtggggccag | | | | | | 249 |

<210> 1466

<211> 203

<212> DNA

<213> Homo sapien

<400> 1466

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| cctcagacac | cttttaattg | cttaggagaa | accattgtct | ctgactgcag | gtttgaataa | 60 |
| gttgaagacc | agagaaaagt | acacactggg | ctacaaagga | atttgagat | agccaaggaa | 120 |
| caggattttcc | cctagcaagc | taccttctgt | tcaaatcatg | aaaaaagact | atttccccct | 180 |
| agaataggga | agcttgctat | ttt | | | | 203 |

<210> 1467

<211> 223

<212> DNA

<213> Homo sapien

<400> 1467

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtcagaac | aggaacgacc | tgggttatgg | aagcccagaa | agggaggagg | acttcttttg | 60 |
| gtcccagtga | aagatgcttc | cagaatctgt | agccttactt | atttgcttgg | atctcactgg | 120 |
| aataacttgg | tggtgaggtc | accggttctg | gggtgatcac | tgggtttgct | gcatagatgt | 180 |

```
<210> 1468
<211> 177
<212> DNA
<213> Homo sapien
```

```
<210> 1469
<211> 185
<212> DNA
<213> Homo sapien
```

```
<210> 1470
<211> 482
<212> DNA
<213> Homo sapien
```

```
<210> 1471
<211> 257
<212> DNA
<213> Homo sapien
```

```
<210> 1472
<211> 342
<212> DNA
```

<213> Homo sapien

<400> 1472

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| cttttgcgag | cctctgccgc | agcagctccg | ttttcacgcg | catctcgttt | ttgtgtgtgt | 60 |
| gtttttgttt | tgtttttgtt | tttgtttttt | tgtttcagag | aattggaagc | taaagctacc | 120 |
| aaagacgtag | aaagaaatct | tagcaggtaa | gatgggagag | ctttccgtct | cccgcaccac | 180 |
| gataatcgta | tattttctact | ccgattcgcc | ctttctgggt | tgagaagttc | ccccgtgaca | 240 |
| ttttcttccg | cacccggaga | gcagacattc | gggagaagcg | gcctggggga | atactggagg | 300 |
| gattgcgggg | agatgcgtaa | ttacgcgtgt | gtttctttct | tt | | 342 |

<210> 1473

<211> 526

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(526)

<223> n = A,T,C or G

<400> 1473

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgctacatg | tcttcacagc | ccaggaattc | aaggcccagg | tggcagcagg | aagaaacagt | 60 |
| ggaaaagcaa | ggggaagaga | aaagagaaaa | aggaggggga | aagtctgcat | aactgtcata | 120 |
| acctctgctt | ctcctgctct | gtaacaaacc | cacaaccagg | aagagtcatt | gtctggaaca | 180 |
| atcatgggac | cccaaacgcc | tgtaggtttt | ttaccacca | acatcaccca | tggtctctct | 240 |
| aagctgtcat | ttgtttccca | cagttacct | gcacacgga | tgcccaattt | atggcccagg | 300 |
| aaggctgacc | caggctaagg | gcagtctcac | tccacagcca | tgcaatggac | agtctgaatg | 360 |
| tttcttacc | cagaccttta | ctgacctcta | ctatttctct | ctctgatata | aaagaaaaac | 420 |
| acttttaatt | ttctnctgca | tnctacatct | cctnctaaaa | antttggcct | aattgncatc | 480 |
| aaaaccttgt | aggaatctga | aatttttggt | cttctgaatc | ttancc | | 526 |

<210> 1474

<211> 187

<212> DNA

<213> Homo sapien

<400> 1474

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| aaacttggtt | gctgtgaaca | attgtcga | agagtcctcc | aattaatgct | ttttatatct | 60 |
| aggctacctg | ttggtttagat | tcaaggcccc | gagctgttac | cattcacaa | aaaagcttaa | 120 |
| acacattgtc | caaaaaaaaa | aaaaaaaaaa | gccccykccc | sgggggscc | ttmaaggggr | 180 |
| aawtccc | | | | | | 187 |

<210> 1475

<211> 474

<212> DNA

<213> Homo sapien

<400> 1475

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccattctctt | tatctcaa | cgaagaaaga | tatgatgcag | gcagtagttt | tttcttagtg | 60 |
| cctcatagta | tctaatagca | gaaagtgcag | cgcatagcgg | agcacattag | tttttatgta | 120 |
| tctacaggac | agaagggcca | cttagctgat | ggctccagg | ttcctttgat | ataatcta | 180 |
| gttcttatga | cctcaaagac | tgaacacatt | tccctaagtg | cttcacttag | caccaggag | 240 |
| caacttggag | tcttcgcaga | ataaaatcca | ttattttaat | gtagattaat | acatgggtac | 300 |
| ttatatctat | gcaggtctat | aatagtttat | tcctatgtaa | gctttattaa | aagcattggt | 360 |

```

atgtttttaca taaaaagtta atgtgaatat tagaaaaaaa ggacaatatt aaagcagttt 420
gtagaatttg tttccccccc aaaatgaatg aaatacacaa tagatgtaca aaaa 474

```

```

<210> 1476
<211> 401
<212> DNA
<213> Homo sapien

```

```

<400> 1476
ccttggggac agggcaggag gacgcacacc tcatggacag ggcggccagg gctgagatac 60
cagcgggggtg ggtattcccc gcgggtgctt acctccaaca gtgtcttgtc agcaaaggcc 120
atgatgcctt caaagatgat gacgtttgca ccatacagt tttctgtga agaaaccag 180
gagttgcgga gcctggctca tgtgcctgca gcccccgag gccccctctg cagggccctg 240
gcctaccag tccttcttcc ggctgtgcgt ggtgaagtca taaatgggca ccttgacact 300
cttccccctgc ttcagcttct tgaggggtgga aatgatgaag gtcgaagtca aaaggcatct 360
gggggtgggtc gaaagtttga aagtttgctt gtgggtgccgg g 401

```

```

<210> 1477
<211> 753
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(753)
<223> n = A,T,C or G

```

```

<400> 1477
cagcatgctt aaaaagttag aggaattgga acagaaatac acctwmcaac ctkrmcctnt 60
taccaaaaac aaacnagtgg tatkggamcc sacctttmrk ctttttcmac macttatttc 120
aaagytsrnt kgtggkgaaa agmcacycyk snatscywcc rcaccttgw aggcygttgg 180
acttrataac akknctgctn atnwnrtgta ggggtgatay tgatgrtgaa attgcactta 240
gctgggttat aattkgaaag tcaaagtctt atttgataaa gatgtgaatg agagaaatac 300
agtaaaagga ttttaggaagt tcaacatttt gggcacgcac acaaaagtga tgaacatgga 360
ggagtccacc aatggcagtc tggcggctga atttcggcac ctgcaattga aagaacagaa 420
aaatgctggc accagaacga atgagggctc tctcatcgtt actgaagagc ttcactccct 480
tagttttgaa acccaattgt gccagcctgg tttggtaatt gacctcgaga cgacctctct 540
gcccgttgtg gtgatctcca acgtcagcca gctcccgagc ggttgggcct ccataccttg 600
gtacaacatg ctgggtggccg gaaccagga acctgtcctt cttoctgact ccccccttg 660
cacgatgggc tcancttttc anaagtgcct gagttggcag tttttcttnt tgtaacccaa 720
aagaaggtct caatggnngg acccanaacc ttt 753

```

```

<210> 1478
<211> 421
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(421)
<223> n = A,T,C or G

```

```

<400> 1478
aaacctatac tcactttccc aaattgaatc actgctcaca ctgctgatga tttagagtgc 60

```

| | | | | | | |
|------------|------------|-------------|-------------|-------------|-------------|-----|
| tgtccggtgg | agatcccacc | cgaacgtctt | atctaatacat | gaaactccct | agttccttca | 120 |
| tgtaacttcc | ctgaaaaatc | taagtgtttc | ataaatttga | gagtctgtga | cccacttacc | 180 |
| ttgcatctca | caggtagaca | gtatataact | aacaacccaa | gactacatat | tgctactgac | 240 |
| acacacgtta | taatcattta | tcatatataat | acatacatgc | atacactctc | aaagcaaata | 300 |
| atTTTTcact | tcaaaacagt | attgacttgt | ataccttgta | atttgaaata | ttttctttgt | 360 |
| taaaatagaa | tggtatcaat | aaatagacca | ttaaccaana | aaaaaaaaaga | aaaaaaaaaaa | 420 |
| a | | | | | | 421 |

<210> 1479

<211> 214

<212> DNA

<213> Homo sapien

<400> 1479

| | | | | | | |
|------------|-------------|------------|------------|------------|-------------|-----|
| ggaaatatat | aataaaaaatg | ttaaccagaa | ggtaaacttg | agtgtaatg | tcagacagac | 60 |
| acacttttcc | accagtgtat | ttgaatttta | gaccagtgc | cctgttttgt | ggcattcatg | 120 |
| caaaacatgc | tgagggcttt | gttcatctgg | tcacgtgtgc | caaatttcag | tcattgtttgt | 180 |
| agcaagattt | tggaagcatt | catatttcct | tttt | | | 214 |

<210> 1480

<211> 434

<212> DNA

<213> Homo sapien

<400> 1480

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| ggaggccgct | tacgtaaagc | ccaggggaca | ttcaacagcc | cctactacc | aggccactac | 60 |
| ccaccaaca | ttgactgcac | atggaacatt | gaggtgccc | acaaccagca | tgtgaagggtg | 120 |
| cgtttcaaat | tcttctacct | gctggagccc | ggcgtgctg | cgggcacctg | ccccaaggac | 180 |
| tacgtggaga | tcaatgggga | gaaatactgc | ggagagaggt | cccagttcgt | cgtcaccagc | 240 |
| aacagcaaca | agatcacagt | tcgcttccac | tcagatcagt | cctacaccga | caccggcttc | 300 |
| ttagctgaat | acctctccta | cgactccagt | gacccatgcc | cggggcagtt | cacgtgccgc | 360 |
| acggggcggt | gtatccggaa | ggagctgcgc | tgtgatggct | gggccgactg | caccgaccac | 420 |
| agcgatgagc | tcaa | | | | | 434 |

<210> 1481

<211> 131

<212> DNA

<213> Homo sapien

<400> 1481

| | | | | | | |
|------------|-------------|------------|-------------|------------|------------|-----|
| aaaatcccca | taaattctttt | ctgtcctgag | gtagttgcaa | aataaatcat | aacttggata | 60 |
| tcaactagag | ctgaggcttt | gactttttac | tcattaaaaac | tagttgttac | aggaactacc | 120 |
| tttagatatt | t | | | | | 131 |

<210> 1482

<211> 324

<212> DNA

<213> Homo sapien

<400> 1482

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| tgctcgctcc | tcagaggctg | aaaacatgag | aagctagggtg | tggtgaaacc | aaagcagctt | 60 |
| tattgttcaa | atgctaaaga | cgggaggatg | gactggctca | agccttaaag | aaaccatctc | 120 |
| gactttttga | actcagtga | cgggtttaag | gaaaacgtgg | gaaatatgca | aaggtggtgc | 180 |
| aggagggtgc | aggtctgtgt | gtcttattcc | catggatatc | ttgagtaatc | gcttgtccag | 240 |

aggtgggggtt tgtgtcatcc tgaattcaac ccagcaatgg taggggtactg ttcataactc 300
accctaagcc agaagattcc tcag 324

<210> 1483
<211> 393
<212> DNA
<213> Homo sapien

<400> 1483
atgtttaatg aatgatacag gatacatccc tgttgggaagc ttgcaaaaaga cacatacact 60
gtgggtacata tttgatttaa tagaagttgt ttatcaggct atatatatat ttgccccaaac 120
atgcaccaca ggataaaata actattttaca taacataggg tatttaattg acatagacta 180
tcagctttgc tgagagcaga agatggcaaa gcaatactgc agcagaaagt ggaacaacta 240
ttctaaagca atacttttaga tatatttttc tagaatggat ttattagatt actttttgga 300
aagcatttga cctaaattaa atatagagct ctgaaactta gaataaaaatt tgcacttgct 360
gaaacagaat actttgcata aaaataatcc ttt 393

<210> 1484
<211> 323
<212> DNA
<213> Homo sapien

<400> 1484
tttagatcag aaagtttgag gtcttcatca gcagacactc gtgcttctat ttttcttggt 60
ttatcgaaca gttctgaaac tttgagaaaa aacttgcata tatctgtaga atcctgagtt 120
cctaaagcat ataatgaaga accaattcta ttgtaatcat ctgcagcact tttgtgggat 180
cttgtcattc tatcagattt agcagatgca tccttaactc ggttatgata ttccaaaaga 240
aatgttcggt cgtgctcaaa gaaatcatct acatccttta ctctgaaac gattactcca 300
tctgctgatt taaccatggt ttt 323

<210> 1485
<211> 405
<212> DNA
<213> Homo sapien

<400> 1485
aggagcgtca ggaaaacacg ggcagcctgg gctctgaccc gagccactcc aactccacgg 60
ccacgcagga agaagacgag gaggaggagg agagtttttg gaccctctct gacaaatact 120
cctcccggag actattccgc aaatccgcag cccagttcca taacctgcgg tttggggaac 180
ggagagatga gcaaattgaa ccggagccca aattatggcg aggccggaga aacaccccgt 240
actggtactt cttgcagtgc aaacacctga tcaaggaagg gaagctggtt gaagccctgg 300
acctgtttga gaggcagatg ctgaaggagg agcgattgca gcccattggag agcaactaca 360
cgggtgctgat tgggggctgc gggcgggttg gctacctgaa gaagg 405

<210> 1486
<211> 230
<212> DNA
<213> Homo sapien

<400> 1486
aaaaatatgt ggattgtgct tgacgtagca aatttcttct atctgcaaaa gcccttttct 60
cactacctca tataaccccc tttgatatgg caccatgttt gaaattggag cgtacacaca 120
tagtcattgg atttactggg attctctttg tgacaagtag gagccaaggg gtcattgcagg 180
gaagcgaacg tgcccagataa ggatttcctt gttgccagag tgtttagcag 230

<210> 1487
 <211> 273
 <212> DNA
 <213> Homo sapien

<400> 1487
 tttccactct gcacattgta gagggaacac tctgtaggcc catgggtccc ttactagaga 60
 ggttgagtga atttgccttc agttaacatg ggaccttctg tttagcttcc tcttgcttcc 120
 caaagatttt aagcattttg taaatgtata aactcacctc tggtaacagt ggcccagacg 180
 ctgctttgtg ctaaaagcat gggaaatgta aaggcagtct ttctctggga aatggatgct 240
 attctattct gctgccctta cctgttcctg agg 273

<210> 1488
 <211> 452
 <212> DNA
 <213> Homo sapien

<400> 1488
 cctactgtgc cccgtaggca aagctctgaa gatttcatcg aaaaatctgc tgtcaatacg 60
 tagaaaagtt cactatttca gtttcacagc aaaaaagggtg gggggagggg ggaacccaat 120
 agatatttaa gtagatgctt tccaatccca ttcactgcat taattagctt acctcttata 180
 cagtacaaca taaacattgc atgtttatct gtatgtaaca cctataagca tatagcatct 240
 acatttttaag tgtattttaca aattcaacaa aatatctaca tataaaaagc tttactttaa 300
 attaaacttg atgcaagtta tgagaaacca atttattggc aaatgaaact gagcattcct 360
 tcaaccatag gttgttatag attttcatat ttggaggtaa cccatttgat agatattggt 420
 tatgaatacg atagaatata tatttacttt tt 452

<210> 1489
 <211> 653
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(653)
 <223> n = A,T,C or G

<400> 1489
 cctgctcttc tcttcaaagc acttagtaca cagggktaca ggtgctacca cttggattcc 60
 ccagagcatg gaagtctgat cccaggttga acatatttct tctgaaaatg agcatcttgg 120
 ttctatagat tcttatcttg ctcacaggac ttgctccaaa actgaatttt cagaagcagc 180
 atgataggga aagagatatt caactctgac agacaaggta gatcgaagca cccacactaa 240
 tttctttcag gtgccccatg aggaagactg catcatgtca ctccactca cttggggaga 300
 ttctaggact gagacacaaa gttccccag agtttctgct aatggaaggg gaaacagggtg 360
 gtttggaatg gaaagggtgga accaggtcca caaatgtgc tccctctgct caagactgac 420
 tttggctttc ccagggtccc acttgacttt catataagct gagatgacct attacgggaa 480
 aaattaggga acacctaata aaaccaactt tcaaaaactc ctatttatca tggatgtgcc 540
 acgatcgaga gaatcnaaca cnaactgnct gtnagagagg ccttcattnt gnctcatctt 600
 gagctaaaat cctgrcttgg gatgccagaa ancatgnccc tcttntcggg ttg 653

<210> 1490
 <211> 363
 <212> DNA

<220>

 $\langle 222 \rangle \quad (1) \dots (363)$

<400> 1490

<210> 1491

<212> DNA

<400> 1491

<210> 1492

<211> 184

<212> DNA

<400> 1492

<210> 1493

<211> 273

<212> DNA

<220>

<221> misc feature

$\langle 222 \rangle$ (1) ... (273)

<400> 1493

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| aggttaawttg | tgatatttag | tgcacattta | cgtgtaggnc | crtcttkaat | ggtaaagaca | 60 |
| gatacaagcc | tatggcacac | ttctccaaag | caagctatac | ttgagagcca | attcccaa | 120 |
| aagacagcag | agatctgact | aaatgcaact | gtgcaaacat | tcaacagaca | tgttgaatgt | 180 |
| aagacaaatt | atgattactg | ataaatatgc | aatgtggctc | ataaatttat | gaatgtgact | 240 |
| tccaaagggga | atatggtatg | gaagcccatt | ttt | | | 273 |

<210> 1494
 <211> 343
 <212> DNA
 <213> Homo sapien

<400> 1494
 ttggaaagcc tatcactttc tctcttcatt ctccagcccc cacaccaagc acacagagct 60
 ttccagtgtc ttactcttaa tggagaacat aaccagggat tatcagggtat tccaacatga 120
 aaaagaaagt ccaatagaaa caagcaggat aatcaaacca ggaggaagca gagactatat 180
 agagaaagaa aaaaagacac atgggaataa cggcaataat actgacaata cacctacca 240
 taaacttatc agaatgaatt tgttggagaa atatatggag gggagggtact tgtgtgtgtg 300
 cacaggcact catgtacacg tgtgtatgtg tatgtttttt taa 343

<210> 1495
 <211> 378
 <212> DNA
 <213> Homo sapien

<400> 1495
 tagcattctt ccagccactc tggcgctact atgtgcttca cgacagaaat cgccgtcagg 60
 aacttcacgg tgcgagtcac tttgctggca atgaggtgtg tgcacttctg tgcagactcc 120
 gcaacctctc caccaagaat gtagagcttc ttaataact gttgaacctg gacaggctcg 180
 aatccagtga aaagcacaaa aggggtcaat tctggagtta gcttttttagt gggaggtggt 240
 acgtcttcaa ttctggctct tttggaagaa ggctggacat tagctacttc attctgtttc 300
 agtttgggag gtagtcttat actcatcaac aactctgcag acacttttaa gggaactctc 360
 caagcatcta aaagattt 378

<210> 1496
 <211> 181
 <212> DNA
 <213> Homo sapien

<400> 1496
 tggagaagga agttttcctg aagagccaga atccttgcta agtcatttag atccaactga 60
 ccatctttat ttctgtcaaa aatcttcac atggtgccag tgtattcttc cagtttagcc 120
 tcagaaatgg cctttttgtg gtgaagaaag aggtctcgga ggaagttgag gagctcagca 180
 g 181

<210> 1497
 <211> 373
 <212> DNA
 <213> Homo sapien

<400> 1497
 tggaaagctga tccaccttga gatcaagccg gccatccgga accagatcat ccgcgagctg 60
 caggtcctgc acgaatgcaa ctgcgcgtac atcggtgggt tctacggggc cttctacagt 120
 gacggggaga tcagcatttg catggaacac atggacggcg gctccctgga ccagggtgtg 180
 aaagaggcca agaggattcc cgaggagatc ctggggaaaag tcagcatcgc gggttctccg 240
 ggcttggcgt acctccgaga gaagcaccag atcatgcacc gagatgtgaa gccctccaac 300
 atcctcgtga actctagagg ggagatcaag ctgtgtgact tcggggtgag cggccagctc 360
 atcgactcca tgg 373

<210> 1498
 <211> 337

<212> DNA

<213> Homo sapien

<400> 1498

| | | | | | | |
|------------|------------|------------|------------|-------------|-------------|-----|
| gctctttag | tgcttttctt | ttaagggaga | tgtagtaaaa | gggaaaatgt | agctcttagt | 60 |
| ttacacttca | aagatgtggg | ggtctttcag | agaactaaga | ataacagttt | tatgtgcaga | 120 |
| gagagtttgc | cagatctgaa | gcatatacct | cattgactag | gctgttactt | tgggataggt | 180 |
| tgcagtacca | gccacagcca | gcagatagag | gaaaagacac | acataaaactc | gcttctgagc | 240 |
| gtccacttct | gcactctctg | ctctgctgtt | actcagcccc | tgagtctgac | tcattctctgc | 300 |
| acaacctctc | tgtgccatga | agataagtct | tccatgg | | | 337 |

<210> 1499

<211> 314

<212> DNA

<213> Homo sapien

<400> 1499

| | | | | | | |
|-------------|-------------|------------|------------|------------|------------|-----|
| catgctggagg | gacttttagca | tggctgataa | ggtccttctc | accattccaa | aagaacagag | 60 |
| gaccagagtt | gcacactttt | tggaaaggca | gggcttcaag | cagcaagctc | ttacagtatc | 120 |
| cacagatcct | gagcatcggt | ttgagcttgc | tcttcagctt | ggagagttaa | aaattgcata | 180 |
| ccagtttagca | gtggaagcag | agtcagaaca | gaagtggaaa | caacttgctg | aacttgccat | 240 |
| tagtaaagt | cagtttggcc | tagcccagga | gtgcctgcat | catgcacagg | attatggggg | 300 |
| cctgctgctt | ttgg | | | | | 314 |

<210> 1500

<211> 321

<212> DNA

<213> Homo sapien

<400> 1500

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| cctgaaacct | ggtgggaaga | tgattgaaag | tgttttagat | tcaacagatt | gactatgtat | 60 |
| gacttatcta | ttaaaatgaa | gaacttccat | ggtttaatag | aatgaatgct | gtattcaaca | 120 |
| aggtcttcca | tccttcttat | aaatcttaag | actgtgttta | agctttcttt | cacttttact | 180 |
| ctatcccttg | gaagttaatt | gggaataaaa | agatttatca | atttagtcac | tataatttaa | 240 |
| ggccaggcat | ctgcttgga | atacaataac | cacaattaat | acttagagaa | aattgtttca | 300 |
| acagattaac | tctgctat | t | | | | 321 |

<210> 1501

<211> 557

<212> DNA

<213> Homo sapien

<400> 1501

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ctgctctggg | gaaaatggtg | gaggagccag | gcagagagga | ggagcagagt | gctggcagtg | 60 |
| gaaagcctag | ctgagactgg | agatgcccc | ctgcccagg | catctcagcg | aggatgcttc | 120 |
| tccatattggg | tgagccagcc | tagagacaga | acaggggaag | ccagcgggtg | ctgcagcgac | 180 |
| ccaccgcccc | agaacatctg | catcttacat | caacaaaggt | ttatttctca | ttaatatcca | 240 |
| ttgtgggttg | gctgccactc | taacctctgt | tgctctcca | tctgggtctt | gggtggcaga | 300 |
| gcagcctgtc | tctgtggcag | aggaaaagag | agcactgggc | agcacaggct | gactctcaaa | 360 |
| ttttccgcct | gaaggtgacc | caagtcactg | ctcacatttc | attgactaaa | gcaaaatcct | 420 |
| atgctgtggg | gtgagttgag | caacgtgatg | aggtgttaac | ttcctacagg | gaggggctca | 480 |
| aatattgccc | aacagtggta | tggccactg | cctgggggtg | tcggtggaag | gctggcagga | 540 |
| caaggagagac | cacgtgg | | | | | 557 |

<210> 1502
 <211> 249
 <212> DNA
 <213> Homo sapien

<400> 1502
 cctgcgggga ggcgcgctgc aagaacctgc ccggctccta ctctgcctc tgtgacgagg 60
 gctttgcgta cagctcccag gagaaggctt gccgagatgt ggacgagtgt ctgcagggcc 120
 gctgtgagca ggtctgcgtg aactccccag ggagctacac ctgccactgt gacgggcgtg 180
 ggggcctcaa gctgtcccag gacatggaca cctgtgagga catcttgccg tgcgtgccct 240
 tcagcgtgg 249

<210> 1503
 <211> 302
 <212> DNA
 <213> Homo sapien

<400> 1503
 ccaggacctc ttttgggcat ttcttcctaa gtggaataca caacagataa gggagtaggg 60
 gaggtaatac agggaagcta ctctttccag ctccagaagga gttgatgaag cccatatatg 120
 cattcaagaa gcccatggga tcctctagct gtggatagtg gctaattgtg tcatccagaa 180
 tcgacactgt ggaccgcggc agcgttttcc tgtacagctc caaaaactct ggatagggat 240
 ttacaggatc caatggccca tagataaaat gaatggggat agttacagag gcaagagctc 300
 cc 302

<210> 1504
 <211> 430
 <212> DNA
 <213> Homo sapien

<400> 1504
 ccacgatatc aactatttgg ctttgtcagg tgttctctca aaaattggca gaagtgggtga 60
 gaatccgtat gccccgctga atctctggc tgactttgct ggtgggtggc ttatgtgtgc 120
 actgggcatt ataattggctc tttttgaccg cacacgcact ggcaaggggc aggtcattga 180
 tgcaaatatg gtggaaggaa cagcatattt aagttctttt ctgtggaaaa ctccagaaatt 240
 gagtctgtgg gaagcacctc gaggacagaa catgttggat ggtggagcac ctttctatac 300
 gacttacagg acagcagatg gggaattcat ggctgttggg gcaatagaac cccagttcta 360
 cgagctgctg atcaaaggac ttggactaaa gtctgatgaa cttcccaatc agatgagcat 420
 ggatgattgg 430

<210> 1505
 <211> 164
 <212> DNA
 <213> Homo sapien

<400> 1505
 ccagtcacct tcaccttcta actaactagc ctccggatga ggtggctgcc accaggcccc 60
 aatgatcccc aggagcccag cttccaaacc ccaacatcga atcaaacatc tccatcccca 120
 agtgcagtaa cacacaaaaa ccaaacactc tgccctggga aagg 164

<210> 1506
 <211> 189
 <212> DNA
 <213> Homo sapien

```
<210> 1507
<211> 268
<212> DNA
<213> Homo sapien
```

```
<210> 1508
<211> 159
<212> DNA
<213> Homo sapien
```

```
<210> 1509
<211> 234
<212> DNA
<213> Homo sapien
```

```
<210> 1510
<211> 437
<212> DNA
<213> Homo sapien
```

| | | | | | | | |
|-------------|-------------|-------------|------------|------------|------------|--|-----|
| <400> | 1510 | | | | | | |
| aaagcagtag | atcttaatat | gaagacagga | atttctatga | tgtttacgaa | cattagactc | | 60 |
| aacattttttg | cagccccctt | tcttgggtcta | cattcacaca | aacatgagac | acagtcccaa | | 120 |
| gggagaaaaca | gatgctggag | gagcatttag | ggccagagtg | gaggcacaga | ggaagctggg | | 180 |
| atcttttcaac | tacccctctc | ttggttactc | ctgggattcc | cttaggattt | cacggcacaa | | 240 |
| ccagcgaaga | gtttgtctcag | attcacttcg | gagtagccac | ttcgggacaa | gaattgctct | | 300 |
| gctgtgtttct | tgagttttct | gtagtctctc | agaacttttg | gggtaaaaaa | ttgtttcttc | | 360 |
| aatttatctt | tctcatgac | ggtagtaagt | ttctccagtg | cacactccgc | atcaaaaatg | | 420 |
| taccggtaaa | agcacag | | | | | | 437 |

```
<400> 1511
tgtgaagatg gagtctgagg ggggtgcaga tgactctgct gaggagggggg acctactgga    60
tgatgatgat aatgaagatc ggggggatga ccag                                     94
```

| <400> | 1512 | | | | | | |
|------------|------------|------------|------------|-------------|------------|--|-----|
| aaaaatatgc | attacaactg | gagttttcca | ctgagaataa | gagtttggtt | ttgacctcmc | | 60 |
| ataaatccaa | gggttcctga | aaaaaaagtt | aatataaatt | ctcaataact | atatcattaa | | 120 |
| taccttatgt | atacatagga | gtttatataa | tgcatttaag | taacaaagaa | tgtaacattt | | 180 |
| attagccacc | aagtaattag | gagatagcat | caattatatt | gaaagaagat | gagtttagat | | 240 |
| gcttatagtc | aagggagtta | attgaaattg | aaagctattg | taggttggtta | ctactattat | | 300 |
| tatcaaacct | gaaagttgga | acatgtgaac | ttgatccttt | gcacacataa | aagttcacaa | | 360 |
| agctgctttt | aatttgccct | tgttctgtag | tactgcttgg | tgaatcatgc | actagtttgt | | 420 |
| tgtaaaattc | atgtaaactt | ttatgtatac | aaatgtcaga | tcaagcacag | gttttattaa | | 480 |
| ttatatatat | ttt | | | | | | 493 |

| | | | | | | | |
|------------|------------|-------------|------------|------------|------------|--|-----|
| <400> | 1513 | | | | | | |
| aaatgaggat | tattgatagt | actcttggtt | tttataccat | tcagatcact | gaatttataa | | 60 |
| agtacccatc | tagtacttga | aaaagtaaag | tgttctgcc | gatcttaggt | atagaggacc | | 120 |
| ctaacacagt | atatcccaag | tgcactttct | aatgtttctg | ggtcctgaag | aattaagata | | 180 |
| caaattaatt | ttactccata | aacagactgt | taattatag | agccttaatt | tttttttcat | | 240 |
| agagatttgt | ctaattgcat | ctcaaaaatta | ttctgccctc | cttaatttgg | gaaggtttgt | | 300 |
| gttttctctg | gaatggtaca | tgtcttccat | gtatcttttg | aactggcaat | tgtctattta | | 360 |
| tcttttat | ttttaagtca | gtatggtcta | acactggcat | gttcagagcc | acattatttc | | 420 |
| tagtccaaaa | ttacaagtaa | tcaagggtca | ttatgggtta | ggcattaatg | tttctatctg | | 480 |
| attttqtgca | aaagcttcaa | attaaaaacag | | | | | 510 |

```
<220>
<221> misc_feature
<222> (1)...(511)
<223> n = A,T,C or G
```

<400> 1514
ctggagatca ggaatagaac ctttccaaga tatcataata ttttctttat aggaacactg 60

| | | | | | | |
|------------|-------------|-------------|-------------|-------------|------------|-----|
| agtaatggca | agaatatattt | gagctttttcc | atgggttaaga | gogatagtct | cagaggctgg | 120 |
| agaaaatggt | cattctgctc | agtgatccag | gagtgtgagg | acagtagctt | cctttccacg | 180 |
| tccacaagac | aatgacagat | gtgttttcctt | ccttgccctt | tctagggatc | tttctagggg | 240 |
| tgttgattct | ctcacaatat | ttcaatgtcc | cattttctgtg | tttctttctcc | ctccaggggc | 300 |
| tgatttacga | ttacatgagt | cttgtcacaa | taattttcctc | ccttaacatc | aaggacaagt | 360 |
| tgatcactga | gataagagct | gatagttcca | tttttattca | gtctccactt | ctgcctgaat | 420 |
| tgcccatggt | cagtccatag | agctacttta | gctccagggtg | tgggtcccggc | cnccatcaca | 480 |
| tcaagaactg | gtttcactgg | gccttggatt | a | | | 511 |

<210> 1515

<211> 176

<212> DNA

<213> Homo sapien

<400> 1515

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| aaaggggaag | gkgaractta | aaagtatttcc | caactagatt | atctacacca | atacattgga | 60 |
| actctatatt | ttgctttcat | tttgtcttaa | aaaaatgaaa | tagcaacgct | ctatcagtca | 120 |
| cacagaggac | atgcarattt | agcagtattg | atattatact | ctatcttgtt | ggattt | 176 |

<210> 1516

<211> 309

<212> DNA

<213> Homo sapien

<400> 1516

| | | | | | | |
|------------|------------|------------|-------------|------------|------------|-----|
| ctggggaaaa | cctgtcatta | cctgcccac | ctgttcatcg | accagctcag | caaccgcgtg | 60 |
| aaggacctga | tggtcataaa | ccgctccacc | accgagctgc | ccctcacogt | gtcctacgac | 120 |
| aaggctctac | tggggcggt | gcgcttctgg | atccacatgc | aggacacogt | gtactccctg | 180 |
| cagcagttcg | ggttttcaga | gaaagatgct | gatgagggtga | aaggaatttt | tgtagatacc | 240 |
| aacttatact | tcctggcgct | gaccttcttt | gtcgcagcgt | tccatcttct | ccttgatttc | 300 |
| ctggccttt | | | | | | 309 |

<210> 1517

<211> 182

<212> DNA

<213> Homo sapien

<400> 1517

| | | | | | | |
|------------|--------------|-------------|------------|------------|-------------|-----|
| ccaacatcta | atTTTTTTTtac | TTTTTaaatta | tagctgttgt | gactgatgtg | agatggcatc | 60 |
| ttactgtggg | TTTTgcttgc | atTTTtttat | ttgatgatta | gtaaggatga | gtgtTTTTTtc | 120 |
| atatacttga | gtgtcttctt | ttgagaaaat | atctgttcat | gtccttttgc | TTTTcttgat | 180 |
| tt | | | | | | 182 |

<210> 1518

<211> 548

<212> DNA

<213> Homo sapien

<400> 1518

| | | | | | | |
|-------------|------------|------------|------------|------------|-------------|-----|
| cctgagggag | agggaaaagc | ggataccac | ctgtgtcgct | gtttgcgtgc | caagtccagg | 60 |
| aacagtccat | acagccctgc | tgcacccac | gacgctgtca | caaagcagga | gttcatccga | 120 |
| ggccaagggtg | ttgtcatgag | aatattcggt | aaagtaggga | cgctgacttt | gttcttgggc | 180 |
| agattctctt | cctgtggagt | atccagcctg | tttgcttagt | tttctgttgc | ttctgggggtc | 240 |
| tgatctctat | ctgttttact | gcagtccagt | taccaaagtg | gtataagtaa | aattgaaaga | 300 |

| | | | | | | |
|------------|------------|-------------|------------|------------|------------|-----|
| cagatggcac | cttgaactca | tttggttaagg | gctgtctcac | tctgccagac | caacaaaaac | 300 |
| tgagactgaa | gtcgccagtc | ctgaggaagc | aggcttgccc | ccagtggaaa | cactcatttg | 360 |
| tcttcagtgg | cgtaacccca | gctcag | | | | 386 |

<210> 1523

<211> 178

<212> DNA

<213> Homo sapien

<400> 1523

| | | | | | | |
|------------|------------|-------------|-------------|------------|------------|-----|
| aaaaagccta | tcccatactg | aattgtggga | acctatgaag | tgtctcttaa | tgtcaattaa | 60 |
| aagtaacagt | ggctgcagat | attgattttct | gaaagtacat | gagaatttgt | ctctaactat | 120 |
| ggttgaaaca | acaaaaccaa | atctgaatca | ggtagagggtc | taccagacac | aaactctg | 178 |

<210> 1524

<211> 319

<212> DNA

<213> Homo sapien

<400> 1524

| | | | | | | |
|------------|-------------|------------|------------|-------------|------------|-----|
| wycacagcwg | aaatggggca | ctgaagtgtg | gagscacaka | atgcggggagg | gcagaaccac | 60 |
| agacaggagg | ctgagattga | cctcctgagt | gcaagctggt | ctccccctca | cctcctgcac | 120 |
| cctacgcaga | tgggtgcttac | cataggattg | ccgtaaaaca | gagacacgca | ccagcgagaa | 180 |
| actttagccc | ttagtateccc | atcctcagga | cagaatcact | cttaaacadg | ttgaaatata | 240 |
| tctgcttaga | gcttttctat | gtgtctatat | aatgtatgca | taatatacaa | ttagaagcat | 300 |
| gtgattttat | aacattttt | | | | | 319 |

<210> 1525

<211> 467

<212> DNA

<213> Homo sapien

<400> 1525

| | | | | | | |
|------------|-------------|------------|------------|-------------|------------|-----|
| ccagactaga | cagagatcag | gtcatcaggg | gagcttccga | gcttcagcaa | agcccacagg | 60 |
| tagctctgcg | aactcagaat | gctaccctac | cttccctgca | ggccgctggt | catgtctgga | 120 |
| ctcctggggg | cgctatttaa | tgtttacccc | catctccagt | gccccctcca | aggctgtgca | 180 |
| gtgtcttggg | gctctcaggg | ccaacatcga | agagatgggg | gccacctctt | aacacctggc | 240 |
| aacagtctcc | cctcatcctg | attcctgaca | acagacaaaa | caccgggtttc | tagggtttat | 300 |
| ctgtttgttt | tttgagttga | gggttcctca | gggccttggc | attgctagtg | atgggtccct | 360 |
| ttgctgtgtg | agaacccccct | caaccccttc | ctcctccctc | tggggatgaa | gtgggagtat | 420 |
| ttggctcccc | atttttgaca | aaagggctca | gtgcagggag | gtggagg | | 467 |

<210> 1526

<211> 439

<212> DNA

<213> Homo sapien

<400> 1526

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| aaactgttta | ctggagaaaa | tctctgctca | tgtccattta | ttgttttttt | ctgtactgtg | 60 |
| atgtgtttca | agcttaggaa | aactagtata | ttagagtatg | ttctaggaaa | ttaaaagatc | 120 |
| tggttagagt | aaaaagttct | ttttaagggt | cttaactaat | tttttcacaa | ctaagaaaat | 180 |
| aaatgaagta | ttcttaggct | gaaattcatc | ttattttatc | ataaattaga | ttgtaggggc | 240 |
| agcctacatt | tttgtgtatg | tgtttttatt | tcttaaatga | ttgtgtgagc | ctgggtgacat | 300 |
| tttatggttc | tttgtatcta | aactgttttt | ccaattcaca | tcttttgtcg | tgaagtgata | 360 |

ttatactaga gtactgtttg cattgtaaaa atgctttgct ggtgctctgg cattttgtct 420
 ttatctcatc acctaattt 439

<210> 1527
 <211> 609
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(609)
 <223> n = A,T,C or G

<400> 1527
 ctggagaact tgggctccat taggtgcaat cgttggagta attagcccat cttttacatt 60
 tcttgccaca aaatctcgaa gagctgccat ttcaggttcg gacagtgaat acacatgtcc 120
 actgggaata ctgtgtgctc caggtatcat ttctatgtga gggccaacca ggcggtgac 180
 tgggtagacg tgctcatcta ctggagtgtg cacattctgg acatagtaat acctcactgg 240
 ttggtaaact ctgtatccat ctactggata atagagtggc ggttgtgggtg ctgggtgggtg 300
 gagcgatggt ggtattggag aatacatccg gcagtggtag cggcagtatt cagaatcaaa 360
 gacgatagat cgagtgtctc atgtgatatt gggatcatgt gtgctcagcc agcgaacccc 420
 taggacgaca ggaagaatg gagactgagt cacatcaaat gacagcacct ctcggtgac 480
 tcccagggtca actatcaggc cgtgagtttc gtggacaact gggcccgatg ctatggggcg 540
 cccatcaatt gcttccacaa gtattggacc cgcccgggcg gncgctcgca agggccgaaa 600
 ttccagcac 609

<210> 1528
 <211> 393
 <212> DNA
 <213> Homo sapien

<400> 1528
 tgatgtaatg aattcatatt tattgatata gaaaaatatg atataatcca tctaaaaagc 60
 aagttacaaa acagtgtaca gtgtaccata gtacctatga acacaattag tgaagtaatt 120
 tgcagagcta taatacaaaa tcagaaatta ttttggtaat gaatttatga ttttcctcgt 180
 tttctgattt tttccatgat ctcatatact ttattctcag aaaacaaaag acaaaacccc 240
 acacatacac aaaaataaac gagtaacttc tttacaaccc cagaggctaa gtcagtggga 300
 aaagagggaa atgaatggtt atgagcataa acacagggac aaataaaaga agtttggagc 360
 acagagaaca attcacaat cagaagtcatt ttt 393

<210> 1529
 <211> 143
 <212> DNA
 <213> Homo sapien

<400> 1529
 atccgataga atccagttca atgaccttca gtctttactc tgtgcaactc ttcagaatgt 60
 tcttcggaaa gtgcaacatc aagatgcttt gcagatctct gatgtgggta tggcctccct 120
 gttaaggatg ttccaaagca cag 143

<210> 1530
 <211> 636
 <212> DNA
 <213> Homo sapien

```

<220>
<221> misc_feature
<222> (1)...(636)
<223> n = A,T,C or G

<400> 1530
gtggagaagc ggcttggtcg ggggtggtct cgtgggggtcc tgcctgttta gtcgctttca 60
gggttcttga gccccttcac gaccgtcacc atggaagtgt caccattgca gcctgtaaatt 120
gaaaatatgc aagtcaacaa aataaagaaa aatgaagatg ctaagaaaag actgtctgtt 180
gaaagaatct atcaaaagaa aacacaattg gaacatattt tgctccgcc agacacctac 240
attggttctg tggaattagt gaccagcaa atgtgggttt acgatgaaga tgttggcatt 300
aactataggg aagtcacttt tgttcctggn ttgtacaaaa tctttgatga gattctagtt 360
aatgctgcgg acaacaaaca aagggaccca aaaatgtctt gtattagagt ccaattgatc 420
cggaaaacaa tttaattagt atatggaata atggaaaagg tattcctgtt gttgaacaca 480
aagctgaaaa gatgtatgtc ccmnctctca tatttgaca gtccttaact tctagtaact 540
atgatgatga tgaaaagaaa gggacaggtg gtcsaatagg ctnttgagcc naattgtgta 600
acatatcag tacccaattt actgnnggaa acagcc 636

<210> 1531
<211> 194
<212> DNA
<213> Homo sapien

<400> 1531
aaaaggcaga gcattctttt ttcggcaatt ttgataagca aggtgtagat ttacattttt 60
gtccttgctc ccaacgaaat ggataaacia aaataactta ccatctactc atggaatgtt 120
gttggttag ccagtctgaa ggccacctt aatttttata taactgtctt tagctcttct 180
tttgacaggg cagg 194

<210> 1532
<211> 300
<212> DNA
<213> Homo sapien

<400> 1532
ccatacaagg taattttgac aggttccttg gattaggaca tgggcatctt gggaggccac 60
tactggccta ccacaactgg gcagcaaaac tattacaccc tccggtataa tagttttgg 120
gtttcaatga ctggggaggaa aagggttggg aatttttgc tttgggtccc tcttaacctt 180
gtatttttaa ggtctgggac tcaccaaccc tccccttcca accagagaaa ctactgcag 240
tatctccttg aaagtctggg gacgagtctg tctaagtgct ggtgagaggc acaggaccaa 300

<210> 1533
<211> 521
<212> DNA
<213> Homo sapien

<400> 1533
gttcctttgc accctgtaga tgttctagga tagttgatgc atgttactaa attacgtatg 60
caagtctgtg agtgcgctcg aggggacatc gccaggact gactgagaca cgatgccgag 120
acctcaagcc ctgaggggga gtcccaaaac ccttacagtg aagatgttta ctcatgccc 180
ccacctctgg tccacactag aaagaagctc gcccacctc cacctgtgag atccgtgaat 240
tctcggaatg gcaggggaag ccttgcacta ggttgagag aagcatcctc cacatcctgt 300
gtcagaaaacc ctggtctccg tggcacttgt aactcacctg gctgtcttct ggtctgtgtg 360

```

```
<210> 1534
<211> 181
<212> DNA
<213> Homo sapien
```

```
<210> 1535
<211> 544
<212> DNA
<213> Homo sapien
```

```
<210> 1536
<211> 591
<212> DNA
<213> Homo sapien
```

```
<210> 1537
<211> 341
<212> DNA
<213> Homo sapien
```

<400> 1537
 acttcggggcc tccctctccc tgtgcagacc ggttgaataa atgataaaat tactgtttgt 60
 gtcctctgtg aagtctggat taatggaaaa aaggatttgt gaggctagtc ttaggctgta 120
 gccaatctgg tgtgcttttt gtgtcttcct gtatggttcc atgataagga ggaatacctt 180
 aggatagaat gcaagcctag gaccccataa gcctgttggt caagccaacc agcaaactgg 240
 gcagtaacaa acattgctgc aggtttccat tttgttttac gtccttgga gcttgacctt 300
 gtaaccacgt ggcagtacct tcttttggcc tctgccattt t 341

<210> 1538
 <211> 363
 <212> DNA
 <213> Homo sapien

<400> 1538
 ggacctgact ttgagtccat cagagacaaa gtgagtgaga tgcacataca gtgtttccag 60
 acctgactca gcccatctgt ctgttaggaa actttatgaa gacgcccccc agaattaaac 120
 cctaattcaa atgtctcact ctgaatagag accttctgaa ataactcttg tatagagacc 180
 cagacacgtg ccttttgcct taaaataaaa atatattagcc catgttggtt tatgtatctg 240
 tctttcagtt agttttgaag gcccgcacgg aaaagtgggg cctgtgcacc tgaaaagaaa 300
 tgtgtatgtt atgtggttgt tggctcttcc tactagagtt atcttgataa ttgtgaagag 360
 tgg 363

<210> 1539
 <211> 371
 <212> DNA
 <213> Homo sapien

<400> 1539
 ctgtgggggt ccttccagag aggagctgag atacgcctac ctggaggggc ccctgggcct 60
 ggaggggctc ctcaagtgtga ctgggtgaag tgttttcaga ggaccagggg tgagggtggg 120
 ggcattctcat ccagaccctg ccggcatctg cccagaacc caagggcccc tcttctctcc 180
 ctctcaatg gaaatgctgg agatgtctc agtcaccctc tgagcactca cacatcacc 240
 cttatttgga aattttctc actctaacct tcttctctgc tgcaccttct gccccatccc 300
 caggtctctg cctctctctc tctcttcta ccttttagca ggtaatgact cagttcccac 360
 tgaggagcca g 371

<210> 1540
 <211> 403
 <212> DNA
 <213> Homo sapien

<400> 1540
 ctkgacgtga tggagcaggt gagcagtgc cgtggggctt gccagagggc tgaggaggac 60
 cctctctaac cagctccctg tcccccttct tctgtagctt gagttgaaga agacactgct 120
 ggacaggatg gttcacctgc tgagtcgagg ttatgtactt cctgttgta gttacatccg 180
 aaagtgtctg gagaagctgg acactgacat ttactcatt cgctattttg tcaactgagg 240
 cagcaatgca ccgttggtt catgtttcat actgtttaca ctagcactgc cctttttggc 300
 ttaatttagt tcattttgta cctaactgag aactgtgctt tctgatgtag tgatgacaat 360
 gacagatact cgtttaccaa aaagcacctt ctgcctgcag cag 403

<210> 1541
 <211> 428
 <212> DNA
 <213> Homo sapien

<400> 1541
 taaaacaaaa ctaaagaaga gaaaatatat tctcgtaaat tatctgaact taaaagatgg 60
 aagcctggag atagatttgt gataagccat tgctgagtac atcctagagt tcttgataat 120
 ttcagttggg taaattacaa tagtttgcta tttcctccct cacattttat gttctacagt 180
 atctagctgc ttgggttttc ctgtatacca tggggcttct gtcactctggg ctttactcag 240
 tggcatattc cctctgccta aaactctcct cccctctcca ccttagaagt agcttttctc 300
 agaacgggtt tcccaggggt tcacctaagg tgatagtaca atctacaggg acctgcacat 360
 gaagaccttt gcatacatgc caggaagttg gactttatct ttggaaaaag ggagcctttg 420
 aagggtttt 428

<210> 1542
 <211> 345
 <212> DNA
 <213> Homo sapien

<400> 1542
 awttaaatgc ttagcaagca gcaattccac gatgggtcaaa ttcctaatat gagagaagta 60
 gaaataggaa aaataggtca ccctgatact tatgttttca ttttgcttaa tatacgtttg 120
 tatatttcaa tataacatta atagatatcg tgtcccttca cagttctaaa gtagtaagca 180
 aaatgaatta atttaaccta tgcaattaaa accaatttgg aagaatattg aggtagcaca 240
 ctgttacggg aattagtagt actcagtaat gcagttgaaa gttagtggct cctaattccag 300
 tatgaatcat ggagatgaga gaaatgatta gataaagaga tatatt 345

<210> 1543
 <211> 420
 <212> DNA
 <213> Homo sapien

<400> 1543
 aatattgaat ttctagaagc agtatattgc ttactgcttc ttaattacgt tatagatgag 60
 gtggaaatga taaaaactaa agaagcaaga ttaatcttta acacacattt caggctgttg 120
 taaaagaata aacaatgctt catataaact tctagcaaat gacttcctaa tgaggctcttg 180
 aaacagtctt tagggcacgg aatgtcatca cataattaag cagctttaag cctttattaa 240
 aaggcttaaa gtcgcaaaca atgaaatctg aaacaaactg taccatatta aactttttga 300
 tgatatttca aattcagtaa aagaaaaaaa ggatgggttca gaataacatc acgtattcta 360
 atcctgaaac acataacaaa tgcactctgaa acagcaattc ttaaaaaggt tttgcccttt 420

<210> 1544
 <211> 306
 <212> DNA
 <213> Homo sapien

<400> 1544
 ctggcttcac tctactccc tctctgctcg cagcacgtcg gccgccagct ctttgatgtg 60
 ttcccaggcc cgctgcacat gggcagattc caccgtgcga gaacagatgg caaagcgcag 120
 gacaaacttg tcctgaggt gacatggaac caagtggatt tttttggcac tgtttattct 180
 ttgcagaaga gcttcattca ctttgttggg acccttttagc cgaaagcaga caagccccag 240
 aatgacttcc acacagattt caaagcgggg atcctggcgc accagtgact caaactcatg 300
 ggacag 306

<210> 1545
 <211> 110
 <212> DNA

<213> Homo sapien

<400> 1545

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|-----|
| ctgtctcggg | ccttcatacct | gaagatcagc | gtgtgcgatg | ccgtcctgga | ccacaacccc | 60 |
| ccaggctgta | ccttcacagt | cctgggtgcac | acgagagaag | ccgccactcg | | 110 |

<210> 1546

<211> 239

<212> DNA

<213> Homo sapien

<400> 1546

| | | | | | | |
|------------|-------------|------------|------------|------------|------------|-----|
| aaagaaatat | gacacgggtgt | tggatattct | aagagacttt | tttgaactca | gacttaaata | 60 |
| ttatggatta | agaaaagaat | ggctcctagg | aatgcttggt | gctgaatctg | ctaaactgaa | 120 |
| taatcaggct | cgctttatct | tagagaaaat | agatggcaaa | ataatcattg | aaaataagcc | 180 |
| taagaaagaa | ttaattaaag | ttctgattca | gaggggatat | gattcggatc | ctgtgaagg | 239 |

<210> 1547

<211> 527

<212> DNA

<213> Homo sapien

<400> 1547

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| aaaaattcca | gttgagattt | ttctggttct | ctgtataaag | attgactgga | acataacat | 60 |
| tttgggggtt | atgtttggag | actttggctc | ttattcaaac | cttccatttt | agttggcttc | 120 |
| ttctgacagt | gcttcagcat | ggaagcaagg | agggggcctc | attactgcca | ggtaagggtta | 180 |
| aaaatctagt | ttctctgctg | ggtctccatt | gtcactaaga | aaggaatggc | tctgttattg | 240 |
| ctgggcaggg | ttggctgttc | caactgataa | tcctatgtct | gggagggcta | ggagtgcctc | 300 |
| cttgctgttc | ctcttgttgt | ttccactgac | agtggagtgg | ccttgttact | gctgggtggg | 360 |
| ggttgagagt | tctggctctc | tactaggagg | gacacaacct | cagtgtagag | aggcggggat | 420 |
| accttgttac | tgtcaggcac | aggcggaggt | ccagtctcct | tactccacct | acccaacagg | 480 |
| gtagcttgag | gcacttcatt | attgcctagt | gagagtggaa | gttttagg | | 527 |

<210> 1548

<211> 333

<212> DNA

<213> Homo sapien

<400> 1548

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgtgggcgg | agctagtagg | ggcggggcta | cgtgattgac | acttctctcc | tcagacttca | 60 |
| agggtacca | ctggaccctt | cccctgtctt | gaaccctgag | ccggcaccat | gcacggacgc | 120 |
| ctgaagggtg | agacgtcaga | agagcaggcg | gaggccaaaa | ggctagagcg | agagcagaag | 180 |
| ctgaagctat | accagtcagc | caccagggcc | gtattccaga | agcgccaggc | tggtagctg | 240 |
| gatgagtcg | tgctggaact | gacaagccag | attctgggag | ccaaccctga | ttttgccacc | 300 |
| ctctggaact | gccgacgaga | ggtgctccag | cag | | | 333 |

<210> 1549

<211> 438

<212> DNA

<213> Homo sapien

<400> 1549

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ttgacagtgt | acgctggagc | aggttccagg | gtggggctgc | cctgccgcct | gcctgctggg | 60 |
| gtggggaccc | ggtctttcct | cactgccaag | tggactcctc | ctgggggagg | ccctgacctc | 120 |

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| ctgggtgactg | gagacaatgg | cgactttacc | cttcgactag | aggatgtgag | ccaggcccag | 180 |
| gctgggacct | acacctgcca | tatccatctg | caggaacagc | agctcaatgc | cactgtcaca | 240 |
| ttggcaatca | tcacagtgc | tcccaaattc | tttgggtcac | ctggatccct | ggggaagctg | 300 |
| ctttgtgagg | tgactccagt | atctggacaa | gaacgctttg | tgtggagctc | tctggacacc | 360 |
| ccatcccaga | ggagtttctc | aggaccttgg | ctggaggcac | aggaggccca | gctcctttcc | 420 |
| cagccttggc | aatgccag | | | | | 438 |

<210> 1550

<211> 204

<212> DNA

<213> Homo sapien

<400> 1550

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaaactaagt | tattccaaca | ctaaaagcat | acaacagcat | gccaacagta | atatattatt | 60 |
| ctccaagact | ttacctatgt | aagtgttcaa | aactctgcag | cattaaacaa | cgtgtatgca | 120 |
| aattgttatg | gatacatctt | agaatctaag | aaatcaggca | agtgtctaaa | aggccaacgg | 180 |
| tccaagggat | tacatctgca | gttt | | | | 204 |

<210> 1551

<211> 132

<212> DNA

<213> Homo sapien

<400> 1551

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ccatctgtgg | atttgtctgt | gcacctattg | gctcttctag | ctgactcttc | tggttgggct | 60 |
| tagagtctgc | ctgtttctgc | tagctccgtg | tttagtccac | ttgggtcatt | agctctgcca | 120 |
| agctgagcct | gg | | | | | 132 |

<210> 1552

<211> 433

<212> DNA

<213> Homo sapien

<400> 1552

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| ctgaatagag | gtcaacacag | ttgcgatgtt | gagggatggt | ctccaagcac | cttttggtgg | 60 |
| caatttgaga | acatccagac | aaatccttcc | agcagaatca | atgtttggat | gataaattgg | 120 |
| agtgagaaat | cggatctgag | gaggttcaaa | tgggtacctc | tcaggaatga | taacttctag | 180 |
| cttaaaaaca | cctttctcat | aagggtgtgt | ggctccacct | aatatttgag | ctcgcaggct | 240 |
| atccatttgg | tctttatctt | gccaacatgt | gatgcctggg | ggtggctctg | tggttaacat | 300 |
| gtgcagctct | ctcttcagac | gtgaagctct | ctgcatgata | cccaagtaga | aggaaccaca | 360 |
| cacagttcac | tgctccacac | taagagctgs | ctgggatgca | ctgagctgac | acccttcaca | 420 |
| acgcagcaac | gcg | | | | | 433 |

<210> 1553

<211> 316

<212> DNA

<213> Homo sapien

<400> 1553

| | | | | | | |
|-------------|------------|------------|------------|-------------|------------|-----|
| gagcaaggtc | tgctgagaac | agacccagtc | cctgaggaag | gagaagatgt | tgctgccacg | 60 |
| atcagtgcc | cagagaccct | ctcggaagag | gagcaggaag | agctaagaag | agaacttgca | 120 |
| aaggtagaag | aagaaatcca | gactctgtct | caagtgttag | cagcaaaaaga | gaagcatcta | 180 |
| gcagagatca | agcggaact | tggaatcaat | tctctacagg | aactaaaaca | gaacattgcc | 240 |
| aaaggggtggc | aagacgtgac | agcaacatct | gcttacaaga | agacatctga | aaccttatcc | 300 |

316

<213> Homo sapien

<223> n = A, T, C or G

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaaggaatta | ttctggcagc | acatgtagta | ttcttggatg | atcttgctgc | tcttatttct | 60 |
| ccttttgtgt | gtgtgtgtgt | gtgtgtggct | atgggttttc | atttgtaact | ccatctgctt | 120 |
| argagagtgg | gctctctata | aggaacctg | ctgtaaacct | cattgcagca | aggatgtaga | 180 |
| gagaaatagg | acttaattcc | actaggggct | ctcatctcac | accttaagga | ggagatttct | 240 |
| agaaaaactg | ggccagattt | tctttgytct | ccatcatttt | aatgtggcag | gctgytcagt | 300 |
| tttcttactc | ttacctatgw | gatatttctt | cgtaacgtgt | ccaaaaagaa | aaaagaccca | 360 |
| atcagtgtct | cttgactttg | ttctttgatc | cctcagtttc | ttcttgattt | cagcatgtgt | 420 |
| ccgggttctc | aattttgggt | atgagttagc | aaatttaacc | attgtgtttg | tgccctaccc | 480 |
| aggggactcc | ccagtttctg | acttgaagta | gactganaag | aatccacgag | gngctatttt | 540 |
| gg | | | | | | 542 |

<213> Homo sapien

ctgtctgtgg cttcccatgt ctttctccaa agttatccag agggttgtga ttttgtctgc 60
ttagtatctc atcaacaaag aaatattatt tgctaattaa aaagttaatc ttcattgg 117

<213> Homo sapien

```
ctgtctgcagc cgcagtttct catccggagt gtaccccgtc atgtcgccgc tggtagcaac      60
gcaaaaggac acggcgcacc ctcgaaactac ggactagtta cttaagcgcg c      111
```

<213> Homo sapien

| | | | | | | |
|------------|------------|------------|------------|------------|-------------|-----|
| cgaggactga | tctcttagta | ctaagtgact | ggggatatta | caytarccaa | cattgggttga | 60 |
| tacatacctk | artmatcatw | tgaggaygca | gtgataarsg | satawwmywg | tatsatccya | 120 |
| acaygyacta | rctcaaaaac | tagtgggggc | ggattgatct | cctgtgggac | wkcacatgsc | 180 |
| ctgaaagtga | acatgmtcmt | ratcacctgc | agrgcttgag | atggyccmca | tkgcwgcact | 240 |
| ccgccccyac | aktttttgaw | tcwacwggag | ttagsswgmt | yctwgawtta | kcctttctac | 300 |
| ctgcctccyg | akagrwgcwc | wygastwggg | kgaatssatt | gackkctaag | rttakacttc | 360 |

cactaactct gtacgmtgar ctcttactaa tattcgttac cacgctaaga ggctctgctc 420
caggatctca tcgcgactgg aaggaacctc cagc 454

<210> 1558

<211> 404

<212> DNA

<213> Homo sapien

<400> 1558

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| aaagaagtgc | agttgatatc | taatttacac | agtgaaacta | gtgatagaaa | ataactaatg | 60 |
| aaaaaaaaatc | agagactggg | ttccaattga | ttgacaccta | gatctgtcag | cctctcttaa | 120 |
| agaaagggga | aggagaaaaa | aaatctcatc | atggaaggca | gacaagagtc | cacctgacag | 180 |
| aggtggaatc | tgatggaatc | tgaccccatc | tcatgataaa | cgagaggaaa | cataaatgcc | 240 |
| atctcaaata | ctaaagcgat | gtagtgtagc | atgagtgact | caatgcaa | tcacagagga | 300 |
| aaagaagtta | cggcttagga | agtaggacaa | taaatacaaa | tatttcatct | tatttaattg | 360 |
| tgcattgactt | cagtgaact | accctttgca | atgcaataaa | tttt | | 404 |

<210> 1559

<211> 266

<212> DNA

<213> Homo sapien

<400> 1559

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| aaactatcag | aagagatgag | agggaattga | tctacaatac | tagaatttta | tgtgcagaca | 60 |
| aatccacatc | tggaaatgaa | atcacagtaa | gatattttcg | ggagaccaa | acataaaaat | 120 |
| tgctagaata | aatttgccac | gaacgagtaa | ctagacatta | gaaattgact | acatagatat | 180 |
| agtaatacta | aaagtgtgta | aaacaagcaa | acacaacaca | cacattctca | attctttttt | 240 |
| tttctatcaa | atatcttcaa | cttttt | | | | 266 |

<210> 1560

<211> 142

<212> DNA

<213> Homo sapien

<400> 1560

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| aaaactcagt | atcttctgaa | ccagaggcat | ttctgattag | cccttcccta | cctattttcc | 60 |
| tagtatcact | ctttaatcag | cttggggagg | tggcagcatt | tcattggcctc | cgtagtaact | 120 |
| cacaatgctt | cctgggggat | tt | | | | 142 |

<210> 1561

<211> 381

<212> DNA

<213> Homo sapien

<400> 1561

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|-----|
| aaacactaaa | tgaagcttct | cacaatttct | aattataaac | aaaaggctga | aaacagtatg | 60 |
| ggaaacaaag | tttcaaaaca | aagaaaagtt | gagtaaaagg | tgccccctct | atggctcatc | 120 |
| tgaaagaaac | atcttactca | gagaggcaaa | catttctgat | ctaggagtaa | gtttcccatc | 180 |
| cactttgcaa | ggaccactc | attctgcaga | aagacctaca | agtctttctg | gtctcaattg | 240 |
| caaagtacgt | gaaaatgtgt | atgaaagatc | taaaagctaa | atattagaat | aaggctaatt | 300 |
| gaaatcaaaa | ttgtgtgctg | gtctaaatat | acattctcgg | cttcttcctt | tttagtaagt | 360 |
| atctttatctt | cagatgtatt | t | | | | 381 |

<210> 1562

<211> 368
 <212> DNA
 <213> Homo sapien

<400> 1562
 ggagaaagga gaaccgtaca tgagcattca gcctgctgaa gatccagatg attatgatga 60
 tggcttttca atgaagcata cagccaccgc cggtttccag agaaaccacc gcctcatcag 120
 tgaaattctt agtgagagtg tgggtgccaga cgttcggtca gttgtcacia cagctagaat 180
 gcaggtcctc aaacggcagg tccagtcctt aatggttcat cagcgaaaac tagaagctga 240
 acttcttcaa atagaggaac gacaccagga gaagaagagg aaattcctgg aaagcacaga 300
 ttcatttaac aatgaactta aaagggttggtg cggctctgaaa gtagaagtgg atatggagaa 360
 aattgcag 368

<210> 1563
 <211> 411
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(411)
 <223> n = A,T,C or G

<400> 1563
 accwtrsaac tgcawttatt acctatgcta gntttggata agaamtgkyc wtayatgtga 60
 kagcaagagg gcacyaraws wrcttsaaca ccaawgggcm ktactwtata kawmcgawgg 120
 gcatgctwtm atgaccaact grmtgactgt ttgagaatgg acaargtgct agcgctaaac 180
 ctgtccttct tgaacrtggc ttgactaacg kcwttgatac gttroccttca kkasaataact 240
 attactasac tttgktgctt gattaccgac tgggtgcactc ttgmtctcac ctatgargac 300
 agtgcctttac acaaactcrt akggaaaatt gnntttgtmc tgtganctac tcatcygaga 360
 nctccctaag ggctaacatt ncatgtttcc gtctcactag ctacacgttc t 411

<210> 1564
 <211> 602
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(602)
 <223> n = A,T,C or G

<400> 1564
 ctagtttttaa gatcagagtt cacttttcttt ggactctgcc tatattttct tacctgaact 60
 tttgcaagtt ttcaggtaaa cctcagctca ggactgctat ttagctcctc ttaagaagat 120
 taaaagagaa aaaaaaaggc ccttttataaa atagtataca cttatttttaa gtgaaaagca 180
 gagaattttta tttatagcta attttagcta tctgtaacca agatggatgc aaagaggcta 240
 gtgcctcaga gagaactgta cgggggtttgt gactggaaaa agttacgttc ccatttctaat 300
 taatgccctt tcttatttta aaacaaaacc aaatgatatc taagtagttc tcagcaataa 360
 taataatgac gataatactt cttttccaca tctcattgtc actgacattt aatgggtactg 420
 tatattactt aatttattga agattattat ttatgtctta ttaggacact atgggttataa 480
 actgtgttta agcctacaat cattgatttt tttttgttat gtcacaatca gtatatatttc 540

tttgggggta cctctctgaa tattatgtaa acaatccaaa gaaatgattg tattaannat 600
tt 602

<210> 1565
<211> 473
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(473)
<223> n = A,T,C or G

<400> 1565
ctagtccagt gtggtggaat tcatccaggg ggctacccct ggctctctgt tgccagtggg 60
catcatcgca gtgggtgtct tcctcttcct ggtggctttt gtgggtgtgt gcggggcctg 120
caaggagaac tattgtctta tgatcacgtt tgccatcttt ctgtctctta tcatgttggt 180
ggaggtggcc gcagccattg ctggctatgt gtnagagat aaggatgatg cagagtttaa 240
taacaacttc cggcagcaga tggagaatta cccgaaaaac aaccacactg nttnatcct 300
ggacaggatg caggcagatt ttaagtgtct tggggctgtc aactncacag attgggagaa 360
aatcccttcc atgtngaaga accgagtcct cgactcctgc tgcattaatg ttactgtggg 420
ctgtgggatt aatttcaacg anaaggcgat ccataaggag ggctgtgtgg aga 473

<210> 1566
<211> 53
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(53)
<223> n = A,T,C or G

<400> 1566
ctagttatta atagnaatca attncggngt cattagttca tagcccatat atg 53

<210> 1567
<211> 136
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(136)
<223> n = A,T,C or G

<400> 1567
ttattgattt ttttttttca ctttcccat cacaactcaca cgcacgctca caactttttat 60
ttgccataat gaaccgtcca gccctgtgg ngatctccta tganaacatg cgttttntga 120
taactnaca ccctac 136

<210> 1568
<211> 192
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(192)

<223> n = A,T,C or G

<400> 1568

```
ttgngtctgt gtagannngt tgaccttctt ccatcccttg gtccttcnct tnccttnccg 60
aggcacagag agacagggca gnatccacgt ncccatntg gaggcagana aaagagaaag 120
tgntttatat acggtactta tttaatatcc nttntaatt anaaantnaa acagttaatt 180
taattaaaga gt 192
```

<210> 1569

<211> 575

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(575)

<223> n = A,T,C or G

<400> 1569

```
ctagttctgt cccccagga gacctggttg tgtctgtgtg agtggttgac ctctctccat 60
cccctgggtc ttcccttccc ttcccaggc acagagagac agggcaggat ccacgtgccc 120
attgtggagg cagagaaaag agaaagtgtt ttatatacgg tacttattta atatcccttt 180
ttaattagaa attaaaacag ttaatttaat taaagagtag gggttttttt cagtattctt 240
ggttaatatt taatttcaac tatttatgag atgtatcttt tgctctctct tgctctctta 300
tttgtaccgg tttttgtata taaaattcat gtttccaatc tctctctccc tgatcgngna 360
cagtcactag cttatcttga acagatattt aattttgcta aactcagct ctgccctccc 420
cgatcccttg gctcccagc acacattcct ttgaaataag gtttcaatat acatctacat 480
actatatata tatttggaac cttgnatttg nngtatata tatatatata tgtttatgta 540
tatatngat tctgataaaa tagacattgc tattc 575
```

<210> 1570

<211> 392

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(392)

<223> n = A,T,C or G

<400> 1570

```
ctagtcacgn gtggtggaat tccgccgcca tcatgggtcg catgcatgct cccgggaagg 60
gctgtccca gtcggcttta ccctatcgac gcagcgtccc cacttggttg aagntgacat 120
ctgacgacgt gaaggagcag attacaaac tggccaagaa gggccttact cttcacaga 180
tcggtgtaat cctgagagat tcacatggtg ttgcacaagt acgttttgtg acaggcaata 240
aaattttaag aattcttaag tctaaggagc ttgctcctga tcttctgaa gatctctacc 300
atttaattaa gaaagcagtt gctgttcgaa agcatcttga gaggaacaga aaggataagg 360
atgctaaatt ccgncatgatt ctaatagaga gc 392
```

<210> 1571
 <211> 390
 <212> DNA
 <213> Homo sapiens

```
<400> 1571
gaaggacgtt tgtgttgga gccctggtat ccccggaact cctggatccc acggcctgcc 60
aggcaggac gggagagatg gtgtcaaagg agaccctggc cctccggggc ccatgggtcc 120
acctggagaa atgcatgtc ctctggaaa tgatgggctg cctggagccc ctggtatccc 180
tgagagtggt ggagagaagg gggagcctgg cgagaggggc cctccagggc ttccagctca 240
tctagatgag gagctccaag ccacactcca cgactttaga catcaaatac tgcagacaag 300
gggagccctc agtctgcagg gctccataat gacagtagga gagaaggctt tctccagcaa 360
tgggcagtc atcacttttg atgccattca                               390
```

<210> 1572
 <211> 383
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(383)
 <223> n = A,T,C or G

```
<400> 1572
ctgcagcttc tgctgctgag gccgggattg ctacgactgg gactgaaggt gaaagaggtg 60
gaatccgaag tcctgggact gcgggatgct aaacattgaa agctgggtgt aggcactgca 120
gggagagtgt ggaggtctga cagggttagga atatgtggga gggctgggct aggaatggcc 180
ttggaggctg gcctgtgtgg atatggcacc aattctaccc tgctcctctt ttcttttccc 240
cagactcaga cgatgccttg ctgaagatga ccatcagcca gcaagagttt ggccgactg 300
ggcttcctga cctaagcagt atgactgagg aagagcagat tgcttatgcc atgcagatgt 360
cctgacangg gagcagagtt tgg                               383
```

<210> 1573
 <211> 149
 <212> DNA
 <213> Homo sapiens

```
<400> 1573
cctccagagc ctctctagtg gcagagcagc tcacactccc tccgctggga acgatggctt 60
ctgcctagta cctatccttg tgtttctgat gcagtggtag cattggttca agttctctcc 120
tgctgtggtc agagttgctt cgatgttg                               149
```

<210> 1574
 <211> 143
 <212> DNA
 <213> Homo sapiens

```
<400> 1574
ctgccaggct gaaaagaagc ctgagctccc acaccgcctt cctcacggcc ctctctcggg 60
agtcatttcc actggtggac cacgggcccc cagccctgtg tgggccttgt ctgtctcagc 120
tcaaccacag tctgacacca gag                               143
```

<210> 1575

<211> 112
 <212> DNA
 <213> Homo sapiens

<400> 1575
 ctgcatccac cctctttcag ggggtagagc cactatactt ctcattgtaga tcagccacat 60
 tgtcactgga gactcggatc cagccatcct cccgcacgtg gtagagggtg ac 112

<210> 1576
 <211> 198
 <212> DNA
 <213> Homo sapiens

<400> 1576
 ccagtatgtc cccaggatta tgtttgttga cccatctctg acagtttagag ccgatatcac 60
 tggaagatat tcaaactgtc tctatgctta cgaacctgca gatacagctc tgttgcttga 120
 caacatgaag aaagctctca agttgctgaa gactgaattg taaagaaaaa aaatctccag 180
 gcccttctgt ctgtcagg 198

<210> 1577
 <211> 444
 <212> DNA
 <213> Homo sapiens

<400> 1577
 cctgcctgga gccccagatc accccttctt actacaccac ttctgacgct gtcattttcca 60
 ctgagaccgt cttcattgtg gagatctccc tgacatgcaa gaacagggc cagaacatgg 120
 ctctctatgc tgacgtcggg ggaaaacaat tccctgtcac tcgaggccag gatgtggggc 180
 gtcattcagg gtccctggagc ctggaccaca agagcgccca cgcaggcacc tatgagggtta 240
 gattcttctga cgaggagtcc tacagcctcc tcagggaaggc tcagaggaat aacgaggaca 300
 tttccatcat cccgcctctg ttacagtca gcgtggacca tcggggcact tggaaacgggc 360
 cctgggtgtc cactgagggtg ctggctgcgg cgatcggcct tgtgatctac tacttggcct 420
 tcagtgcgaa gagccacatc cagg 444

<210> 1578
 <211> 294
 <212> DNA
 <213> Homo sapiens

<400> 1578
 ccacaaagcc attgtatgta gcttttagctc agcgcaaaga agagcgccag gctcacctca 60
 ctaaccagta tatgcagaga atggcaagtg tacgagctgt gcccaaccct gtaatcaacc 120
 cctaccagcc agcacctcct tcagggtact tcatggcagc tatcccacag actcagaacc 180
 gtgctgcata ctatctcct agccaaattg ctcaactaag accaagtccc cgctggactg 240
 ctcagggtgc cagacctcat ccattccaaa atatgcccgg tgctatccgc ccag 294

<210> 1579
 <211> 295
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(295)

<223> n = A,T,C or G

<400> 1579

```
ccacaaaagcc attgtatgta gcttttagctc agcgcaaaga agagcgccag gctcacctca 60
ctaaccagta tatgcagaga atggcaagtg tacgagctgt gcccaaccct gtaatcaacc 120
cctaccagcc agcacctcct tcaggttact tcatggcagc tatcccacag actcanaacc 180
nngctgcata ctatcctcct agccaaattg ctcaactaag accaagtccc cgctggactg 240
ctcagggngc cagacctcat ccattccaaa aatatgcccc gtgctatccg cccag      295
```

<210> 1580

<211> 166

<212> DNA

<213> Homo sapiens

<400> 1580

```
cttcttttatt ggggacatgt gggctggaac agcagatttc agctacatat atgaacaaat 60
cctttatttat tattataatt atttttttgc gtgaaagtgt tacatattct ttcacttgta 120
tgtacagaga ggtttttctg aatatattatt ttaagggtta aatcac      166
```

<210> 1581

<211> 449

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(449)

<223> n = A,T,C or G

<400> 1581

```
ctgaggcaac agaataaatg cagaggcatt acaatgaatc ccaacttaata taaagaacta 60
tacagaccaa cacttctcta caaaatTTTT ttttcctcat tgccagttaa atacagagtt 120
ttactttcat agcttaacaa tgaagggtca taaactgaag ccaatacata tacctagcat 180
ttcagtctaa gcttgtccac gtacatagct gaagtcaatt acaaggtttg gcctagaaat 240
gctaggggaa cttcttttga gttttttacag gtattaaact tcatcttgca cactgaagtc 300
atcatacata cagggcaaaa tcagagcttt tatatttgcg tttattcttc atttaacttt 360
ttataaact actatagttt attaaaacaa aaaacaaaga gcaagtagtg agcatattan 420
gattacagtc ctttctactca ttcacacct      449
```

<210> 1582

<211> 302

<212> DNA

<213> Homo sapiens

<400> 1582

```
ccaatgggct ttgctgtagc ttgctgaaat caccaagcag gagagattta accagaggcg 60
atgtgtccag tcaccagcat agagccatcc tctgtgtcac catccacacg cagggccttc 120
tggcagacct catgcaatgc cctccatgtt aatattcatc agaaaatgga taattagggg 180
ggccagcaaa aatatcaagg gtcaaatac gcacatttct gtttaggcca tctatggctt 240
tcatctcttc tgaagtcaac tggaattcaa acacctgcac gttctgtctg atgcgctgct 300
ca      302
```

<210> 1583

<211> 170

```
ccaatgtaca tgggtggacta tgccggcctg aacgtgcagc tcccgggacc tcttaattac 60
taaacctcag tactgaatca ggacc                                     85
```

<210> 1588
 <211> 369
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(369)
 <223> n = A,T,C or G

<400> 1588
 ccaggctacc ttcccactgg agacaggcag ggggacaggt gctaagggac ctggcaggca 60
 gggctggcag gccccatggc gcctgttcca gcagatgaca agcccaggtc agggtagagc 120
 gggcaggagg ggggacgagg gctcccacaa catgattttg tgtaaaatat ggcagcgaca 180
 cacgctcagg gccgggaggt ggggggttagg gtggggacgg cggcaacatc gtgtaaaaaa 240
 gtgtcccagt tcccatagca aagagagctg tgaccgggtg ttcagagctt ctccagtaca 300
 agggggaaag ccgcccggcg ggggcggcgg gcagggacat catttggttt cctggtgctg 360
 tcngtccga 369

<210> 1589
 <211> 361
 <212> DNA
 <213> Homo sapiens

<400> 1589
 ctgtagcttc tgtgggactt ccactgctca ggcgtcaggc tcagatagct gctggccgcg 60
 tacttgttgt tgctttgttt ggaggggtgtg gtggtctcca ctccgcctt gacggggctg 120
 ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
 agtgtggcct tgttggcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg 240
 gcagccttgg gctgaccag gacggtcagc ttggtccctc cgccgaacag taaaagggga 300
 ctgaggtgtg tatcatagga ctggcagtaa taatcagcct catcttcagc ctggagccca 360
 g 361

<210> 1590
 <211> 434
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(434)
 <223> n = A,T,C or G

<400> 1590
 ctggagaagg tgtgcagggg aaaccctgct gatgtcacgg aggccagggt gtotttctac 60
 tcgggacact ctctcttttg gatgtactgc atggtgttct tgggtgctgta tgtgcaggca 120
 cgactctgtt ggaagtgggc acggctgctg cgaccacag tccagttctt cctggtggcc 180
 tttgcctctt acgtgggcta caccgcgtg tctgattaca aacaccactg gagcgatgtc 240
 cttgttggcc tctgcaggg ggcactggtg gctgccctca ctgtctgcta catctcagac 300
 ttcttcaaag cccgaccccc acagactgt ctgaaggagg aggagctgga acggaagccc 360
 agcctgtcac tgacgttgac cctgggcgag gctgacnaca accactatgg ataccgcac 420
 tcctctcctt gagg 434

<210> 1591
 <211> 439
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(439)
 <223> n = A,T,C or G

<400> 1591
 gctttcgcca gaaaatgttg catgtcaaac aatatgtgat ccatactgtg tgcgtcctt 60
 gggggtttat ttgactttgt cacaatgaca gccaacagtg agactgataa gcctgtaaaa 120
 ataaaaaaat aagactaatc aaatagacat ggcattttta tctcaaagtg caaaatcatc 180
 taactgaaaa tgacggcatt gagaaattcc agtgggttaa aatgaatcaa aacttcatta 240
 cgcaggcagt ggaagtgtgt tgaaagattt accaggggtg tcaagtttta gacactcaga 300
 aaggcaccat tctagccatc ttgattggat aacatgtata tacttatgtc cctacgatat 360
 tcaaaagata atactgtttt agtacaaaac aatcaaaca ggcaaagant caaaaccaag 420
 ccaaccctaa tatccccag 439

<210> 1592
 <211> 74
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(74)
 <223> n = A,T,C or G

<400> 1592
 tttttttttt taatgttcac agtccctgct ttatttccat ttgttcacac acncttttaa 60
 aaaaaaaaaa aaaa 74

<210> 1593
 <211> 288
 <212> DNA
 <213> Homo sapiens

<400> 1593
 ccattccgaag caagattgca gatggcagtg tgaagagaga agacatattc tacacttcaa 60
 agcttttggtg caattcccat cgaccagagt tgggtccgacc agccttggaa aggtcactga 120
 aaaatcttca attggattat gttgacctct accttattca tttccagtg tctgtaaagc 180
 caggtgagga agtgatccca aaagatgaaa atggaaaaat actatttgac acagtggatc 240
 tctgtgccac gtgggaggcc gtggagaagt gtaaagatgc aggattgg 288

<210> 1594
 <211> 455
 <212> DNA
 <213> Homo sapiens

<400> 1594
 ccacacagac tcaccaagcc acagacttgt cttccacaag cacgttctta ccttagccac 60
 gaagtgaacca agccacacgt actaaagggt gaactcaaa atagtgtacag ggtattaaac 120

```
<210> 1595
<211> 367
<212> DNA
<213> Homo sapiens
```

| <400> 1595 | | | | | | |
|------------|------------|-------------|------------|-------------|------------|-----|
| ccaggctacc | tcccactgg | agacaggcag | ggggacaggt | gctaaggggac | ctggcaggca | 60 |
| gggctggcag | gccccatggc | gcctgtttcca | gcagatgaca | agcccagggtc | agggtagagc | 120 |
| gggcaggagg | ggggacgagg | gctcccacaa | catgattttg | tgtaaaatat | ggcagcgaca | 180 |
| cacgctcagg | gcggggaggt | ggggggttagg | gtggggacgg | cggcaacatc | gtgtaaaaaa | 240 |
| gtgtcccagt | tcccatagca | aagagagctg | tgaccgggtg | ttcgagcttc | tccagtacaa | 300 |
| gggggaaagc | cgcccggcgg | gggcggcggg | cagggacatc | atttggtttc | ctggtgctgn | 360 |
| cagtccg | | | | | | 367 |

```
<400> 1596
ctgttcttca tgcgcctggt gggaagacg cccattgaga cactgatcag agacatgctg 60
ctgtcgggga gtaccttcaa ctggccctac ggctcggggc agtgaccatg acgggggccac 120
gtgtgctgtg gccaggcctg cagacagacc tcaagggaca gggaatgctg agggcccggg 180
tgcctctcg agg                                     193
```

```
<400> 1597
ccatgctgga tgttctgctg cttagacctg atctgctgcc aattaccagg ggcagggtcaa 60
ggatgacctt cttggatcca ggaacgctaa catagatcag taaggaatat tcaactcgaa 120
ggatgttgca gccaggata gaagg                                     145
```

<400> 1598
ctgcctataa aactagactt ctgacgctgg gctccagett cattotcaca ggatcatcatc 60

```
ctcatccggg agagcagttg tctgagcaac ctctaagtcg tgctcatact gtgctgccaa 120
agctgggtcc atgacaactt ctggtggggc gagagcaggc atggcaacaa atcccaagtt 180
agggctcca atgagcttcc tagcaagcca gaggaagggc ttttcaaagt tgtagttact 240
tttggcagaa atgtcgtagt actgaagatt cttctttcgg tggaagacaa tggatttcgc 300
cttcactttc ctgtccttaa tatccacttt gttgccacac aacacaatgg ggatgttttc 360
acacactcgt accagatctc tatgccagtt aggcacattc ttgtaagtaa ctctcgatgt 420
tacatcaaac attatgatgg cacac 445
```

<210> 1599

<211> 142

<212> DNA

<213> Homo sapiens

<400> 1599

```
cctgccccag ggggaagcac ggacccgaga cgacggcgat gaggaagggc tcctgacaca 60
cagcgaggaa gagctggaac acagccagga cacagacgag gatgatgggg ccttgacagta 120
agcagcctga caggagcaat gg 142
```

<210> 1600

<211> 297

<212> DNA

<213> Homo sapiens

<400> 1600

```
cctgcacttg aacatggctt tggttttaag caacttctct accctgaccc tcctcctggg 60
acagcgtttc gggaggtttc ttggcctcac tgagagggat gtggagctgc tgtaccccg 120
caaggagaag gtattctaca gcctgatgag ggagagcggc tacatgcaca tccagtgcac 180
caagcctgac accgtaggct ctgctctgaa tgactctcct gtgggtctgg ctgcctatat 240
tctagagaag ttttccacct ggaccaatac ggaattccga tacctggagg atggagg 297
```

<210> 1601

<211> 289

<212> DNA

<213> Homo sapiens

<400> 1601

```
ctggagatga tcctcaacaa gccagggctc aagtacaagc ctgtctgcaa ccaggtggaa 60
tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattggt 120
ctggttgctt atagtgtctt gggatcccac cgagaagaac catgggtgga cccgaactcc 180
ccggtgctct tggaggaccc agtcctttgt gcctcggcaa aaaagcacia gcgaacccca 240
gccctgattg ccttgcgcta ccagctacag cgtgggggtg tggctcctgg 289
```

<210> 1602

<211> 398

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(398)

<223> n = A,T,C or G

<400> 1602

```
gggagggcag agggagaatg ggaagatcag gaagctctag attacttcag tgataaagag 60
```


<400> 1606

```

ccggagccca cgggtggtcat ggctgccaga gcgctctgca tgctggggct ggtcctggcc 60
ttgctgtcct ccagctctgc tgaggagtac gtgggcctgt ctgcaaacca gtgtgccgtg 120
ccagccaagg acaggggtgga ctgcggttac ccccatgtca cccccaagga gtgcaacaac 180
cggggctgct gctttgactc caggatccct ggagtgcctt ggtgtttcaa gcccctgcag 240
gaagcagaat gcaccttctg aggcacctcc ag 272

```

<210> 1607

<211> 444

<212> DNA

<213> Homo sapiens

<400> 1607

```

ccaggctggt ctcaaactcc tcacctcaac tgatccgccc accttggcct cccaaagtgc 60
tggtattata ggtgtgagcc accgtgccca aagttaagta tttttgatca agtgttttgt 120
cttttgtgca aggcatthgt ggctctgtca tagcagagga aaacaaaaca tgcctatcaa 180
atgaatcaag tccgacctct tctcatattg agcaactaga ggtctaggaa catttccct 240
acctgtcatt ctcatctggc ataccagggtg tacatactcc ttcttattct cctctgttac 300
caagatgttg gccccattgg gtttgagggtc acgaacttca caaactccaa actcttggac 360
ctcagtgttg aaggtgaggt catagcctag tgtggagaca tcattttcca gcagataaac 420
cagaccttgg tagaagtggg aatc 444

```

<210> 1608

<211> 189

<212> DNA

<213> Homo sapiens

<400> 1608

```

caaaatccaa aacttctctt gaaaagtcca gggaccgtcc aggggagatg gggaggagat 60
atggagttag tcacctgtct cagaagatgc cagcttctct ctccaggggtg cttagtgtggc 120
tttgcccacc cctcactccc cagggagctc tggggacagc ttctctgcac cctgttccca 180
cccacacag 189

```

<210> 1609

<211> 426

<212> DNA

<213> Homo sapiens

<400> 1609

```

cttttgttat ccttagagga ctcaactggtt tcttttcata agcaaaaagt acctcttctt 60
aaagtgcact ttgcagacgt ttcactcctt ttccaataag cttgagttag gagcttttac 120
cttgtagcag agcagtatta acacctagtt ggttcacctg gaaaacagag aggttgaccg 180
tggggtcac catgcggatg cgggtcacac ggaatgctgg agagatgtta tgtaatatgc 240
tgaggtggcg acctcagtgg agaaatgtaa agactgaatt gaattttaag ctaatgtgaa 300
atcagagaat gttgtaataa gtaaagtgcct taagagtatt taaaatatgc ttccacattt 360
caaaatataa aatgtaacat gacaagagat tttgcgtttg acatttgtgtc tgggaaggaa 420
gggcca 426

```

<210> 1610

<211> 447

<212> DNA

<213> Homo sapiens

<400> 1610

```

cagggctata gtgcgctatg ttgatctggt gttcatgcta agttccgcat caatatgggtg 60
acttcttggg agtggggggac caccagggttg cctaaggagg ggtgaacctg cctacgttgg 120
aaatagagct ggtcaaaaact cctgtgctca tcagtagtag aattgcacct gtgaatagcc 180
accgccctcc agcatgggca acatagcaag accctgcctc ttaagataaa aattggaaaa 240
cactggtagg aaaaaaaggc tgtttgggtc aaataagtct ggattgggta taaatgacac 300
aaaactatca tgaatttgaa agcattttcta atttcttgaa agtctgaaaa agtttaaaca 360
gaatttttagc tgaaaagtcc tgaaagacat ttgaaaaaaa acagcaagaa cacttaaaac 420
tattcaaggt ttgggctggg cacagtg 447

```

<210> 1611

<211> 238

<212> DNA

<213> Homo sapiens

<400> 1611

```

ccaccgggggt tgacctctct cgetagcagg gcccaccag ctcactcccc gcgtcttcca 60
tccccctctag gattccccatt gtccccctact ccagcactag gcaggcaccc ccagcccact 120
gcgactccca ccacgaagga ccccagccct ctctcagcca acacggcccc gccaccgtc 180
tcagacatcg tgcttcttct ggtggggccag gagtctctcc tcgtcgtcga aggtctgg 238

```

<210> 1612

<211> 293

<212> DNA

<213> Homo sapiens

<400> 1612

```

ctgctgcttg tatcctcggg agaggggttc ccactctgag cgggtgggaa ggcaatgcca 60
aacatccggg aaaaataaaa ccactgtctc cacatgagct ggaactgtac gcccttgtg 120
ggtctcctca gggcgatggg agcgaatctc tgcaaaacgg taccattgtg tgcacacact 180
tagatcaatg cctgtcagag cttacaaca acgaatagca gtcttaatca acacagaggg 240
atctttttct gggctctggc catccaacga aggagaccag tggcccccaa tgg 293

```

<210> 1613

<211> 224

<212> DNA

<213> Homo sapiens

<400> 1613

```

ctggattgac cccaaccaag gctgcaacct ggatgccatc aaagtcttct gcaacatgga 60
gactggtgag acctgcgtgt accccactca gccagtggtg gccagaaga actggtacat 120
cagcaagaac cccaaggaca agaggcatgt ctggttcggc gagagcatga ccgatggatt 180
ccagttcgag tatggcggcc agggctccga ctctgccgat gtgg 224

```

<210> 1614

<211> 439

<212> DNA

<213> Homo sapiens

<400> 1614

```

ctccaccctg gcgatggctc cctggtccta ctttctctct caaactgggt ttttctcatt 60
cctttgactc cgccagactt cctcgcccc atgacctggt gttgtgtctg atcacccaa 120
cattcctggc tgcccaatgt ggggcaatga agacccagtg gaaggaatgc tagagtgtgt 180
gaaagtggag gacgcatcgt caaaggacac ctgaggacgt ctcaaagaag ctcggcggga 240
gagctgagcg ctcggaagaa ccaagaatca tctcttttga aaaatcgatt catcaaatga 300

```

```

atcttcggcc aacaactggt caagaaggat tcaaatatca caggttccaa gaagtaaagc 360
tttgagggtc acaaaattag caatagaagc tgggttcgcg catatagatt ctgctcattt 420
atacaaataa tgaggagca                                     439

```

<210> 1615

<211> 237

<212> DNA

<213> Homo sapiens

<400> 1615

```

aggcactcct ggaagtgggt cagtcagggt gcaaaaacat tgaacttgct gtcatgaggc 60
gagatcaatc cctcaagatt ttaaatacctg aagaaattga gaagtatggt gctgaaattg 120
aaaaagaaaa agaagaaaac gaaaagaaga aacaaaagaa agcatcatga tgaataaaat 180
gtctttgctt gtaattttta aattcatatc aatcatggat gagtctcgat gtgtagg 237

```

<210> 1616

<211> 266

<212> DNA

<213> Homo sapiens

<400> 1616

```

ctgggctcta gtttcattcc atctgtcatt ctcaggtaac agggacacat gtccaagtgt 60
tggtcccccgt ggcattgatt tagctttggt gataggcatt gcatcttttg tgtaatatgc 120
aataatggca tgaccagatt catgatatgc tgtgatgggt ttgtttttgt tatcaatttc 180
cacacttctt ctttcaggcc ccattagaat tttgtctttg gaaaactcca gtccttccat 240
ggtaaccatt tcttttccat caacag                                     266

```

<210> 1617

<211> 185

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(185)

<223> n = A,T,C or G

<400> 1617

```

ccatggctag gtttatagat agttgggtgg ttggtgtaaa tgagtgaggc aggagtcgga 60
gnaggtagt tgtggcaata aaaatgatta aggatactag tataagagat caggttcgtc 120
cttttagtggt gtgtatgggt atcattttgt ttgagggttag tttgattagt cattgttggg 180
tggtg                                     185

```

<210> 1618

<211> 354

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(354)

<223> n = A,T,C or G

<400> 1618

```

ctgttaacag ataagtttaa cttgcatctg cagtattgca tgttagggat aagtgcttat 60
ttttaagagc tgtggagttc ttaaatatca accatggcac tttctcctga ccccttcctt 120
aggggatttc aggattgaga aatttttcca tcgagccttt ttaaaattgt aggacttggt 180
cctgtgggct tcagtgatgg ngatagtaca catntcactc agagngcatn tntgcatctt 240
ntaanatana tttcttaaaa gcctctaaag tgatcagntg ccttgatgcc aactaaggaa 300
atttgtttag cattgaatct ctgaaggctc tatgaaagga atagcatgat gtgc 354

```

<210> 1619

<211> 170

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(170)

<223> n = A,T,C or G

<400> 1619

```

ctgtgctgtg gagagaagct gatgttttgg tgtattgtca gccatcgtec tgggactcgg 60
agactatggc ctgcgcctccc caccctcctc ttggaattac aagccctggg gtttgaagct 120
gactttatag ctgcaagtgt atctnncttt tatctggtgc ctctcaaac 170

```

<210> 1620

<211> 386

<212> DNA

<213> Homo sapiens

<400> 1620

```

cctgttgatt gcatactgta gaagatttga tgttcagact gggtcttctt acatatacta 60
tgtttcgtct acagttggta aatttttgtt tttctttgta ttaaattgtg aattgtattg 120
tctggaggaa aagacagagg tctaaaaata aagaaggagt acagtttggg catggtgggt 180
cacccttgga gtcctagcac tttggggggc aaggcaggca gattgcttga gcccaggagt 240
tctagatgag cctgggcaac atagtggagc cccatctcta aaaaaacagt tttagggcca 300
ggcacagtgg ctcacacctg taagcccagc actttggggg gccgaggcag gcagatcata 360
agggcaagag attgagacca tcctgg 386

```

<210> 1621

<211> 346

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(346)

<223> n = A,T,C or G

<400> 1621

```

ccaattctgc ccgttccccg tgggccaaca acaactggggt tgtatgcgtc tggaaacctg 60
tgatagtctt cggcttgcca gcctggccca ccacatccac tgcttgccc acacggacag 120
acactggcaa tggccgcagc tcctcatcaa acgtaaccag cattcggggc tgcattggcag 180
ccaccagccc atacaatata tagtgtgatt tgcctagaat aatgtttcga acatccagga 240
aagagacaag cacagtgagc agtccancca cggccacctg gtcataagc tgccgggtcgc 300
tgtggtaggg gcagagggta aggggtgcct tccctaaatg tgtcag 346

```

<210> 1622
 <211> 366
 <212> DNA
 <213> Homo sapiens

<400> 1622
 ggaagtttgt gctctctgcg tggctaagtt tttcacctac taggacgggg gtgggggtggg 60
 gagaacaggt gtccttctaa aatacagcac aagctacagc ctgctgccag ccataaccca 120
 ggagtaacat cagaaacagg tgagaatgac cactttaact caccggggccc gtcgcactga 180
 aataagcaag aactctgaaa agaagatgga aagtgaggaa gacagtaatt gggagaaaag 240
 tccagacaat gaagattctg gagactctaa ggatatccgc cttactctta tggaagaagt 300
 attgcttctg ggactaaaag ataaagaggg gtacacatct ttctggaatg actgcatatc 360
 atcagg 366

<210> 1623
 <211> 165
 <212> DNA
 <213> Homo sapiens

<400> 1623
 ctgttgattg gctgtgacac tgctttgtgt catcttctta ccatgatcaa aggcgaagga 60
 agggatctct tttgggacat tgtgattggt ttagcagaga gagaaagaga tgaaatacac 120
 ttcggttttc tcttaaaaga tgcattgtatc atacagtgtc ttaag 165

<210> 1624
 <211> 227
 <212> DNA
 <213> Homo sapiens

<400> 1624
 ccaatgcccc gagcaggccc tctttccatc cctgtcggga tgagctgggc aactatgtca 60
 acaaacggaa taccacgtgg caagccgggc acaacttcta caacgtggac atgagctact 120
 tgaagaggct atgtggtacc ttcctgggtg ggcccaagcc accccagaga gttatgttta 180
 ccgaggacct gaagctgcct gcaagcttcg atgcacggga acaatgg 227

<210> 1625
 <211> 373
 <212> DNA
 <213> Homo sapiens

<400> 1625
 ctgtagcttt tgtgggactt ccaactgctca ggcgtcaggc tcaggtagct gctggccgcg 60
 tacttgttgt tgctttgttt ggagggtgtg gtgggtctcca ctcccgcctt gacggggctg 120
 ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
 agtgtggcct tgttgcttg aagctcctca gaggaggggtg ggaacagagt gaccgagggg 240
 gcagccttgg gctgacctag gacggctcagt ttgggtccctc cgccgaacac ccgaagataa 300
 ttagtgctgt ctgttgagta acaatagtag tcaccttcac cttccacctg ggccccagtg 360
 atgggtcaagg tgg 373

<210> 1626
 <211> 367
 <212> DNA
 <213> Homo sapiens

<400> 1626

```

ccagacgtgg tggctcacac ctgcaatccc agcaccttag gaggccgagg caggaggatc 60
cttgaggtca ggagttcgag accagcctcg ccaacatggg gaaaccccat ttctactaaa 120
aatacaaaaa ttagccaagt gtggtggcat atgcctgtaa tcccaactac tcagaaggcc 180
gaggcaggag aattacttga acgcaggaga atcactgcag ccctggaggc agaggttgca 240
gtgagccgag attgcaccac tgtactccag cctgggtgac agagcaagac tccatctcag 300
taaataaata aataaataaa aagcgctgca gtagctgtgg cctcacctg aagtcagcgg 360
gcccagg                                           367

```

<210> 1627

<211> 424

<212> DNA

<213> Homo sapiens

<400> 1627

```

ctggataagg acatcaatac cttctctatg cgtgtcaggg tgtggtacgg gtatcacttt 60
ccggagctgg tgaagatcat caacgacaat gccacatact gccgtcttgc ccagttttatt 120
ggaaaccgaa ggaactgaa tgaggacaag ctggagaagc tggaggagct gacaatggat 180
ggggccaagg ctaaggctat tctggatgcc tcacggctct ccattgggcat ggacatatct 240
gccattgact tgataaacat cgagagcttc tccagtcgtg tgggtgtcttt atctgaatac 300
cgccagagcc tacacactta cctgcgctcc aagatgagcc aagtagcccc cagcctgtca 360
gccctaattg ggaagcgggt aggtgcacgt ctcatcgcac atgctggcag cctcaccaac 420
ctgg                                           424

```

<210> 1628

<211> 314

<212> DNA

<213> Homo sapiens

<400> 1628

```

tcgactgtta tagcttagaa agcaacacta ctactatgag actataaaac attaaactat 60
tttaagaaaa ccacgtgtg gaaaaatgga gccatttttg tcaaaaagtg gctcaaagca 120
caaaactgct cagatgttca agagtcctag gagtctgggc tgcacagtat taaggggtga 180
gaggagaccg acagcctgtt tgaatcaggc ttgtgagccc agctcatctg acaacttcaa 240
agagcttctc tgcctataca ttccaccgtt tagcataaga caccacttta cgctattttac 300
aagtctcctt ttgg                                           314

```

<210> 1629

<211> 393

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(393)

<223> n = A,T,C or G

<400> 1629

```

ctggaccagc accccattga cgggtacctc tcccacacog agctggctcc actgcgtgct 60
cccctcatcc ccattggagc ttgcaccacc cgcttttttc agacctgtga cctggacaat 120
gacaagtaca tcgccttgga tgagtgggcc ggctgcttcg gcatcaagca gaaggatatc 180
gacaaggatc ttgtgatcta aatccactcc ttccacagta ccggattctc tctttaaccc 240
tccccttcgt gttttccccc aatgttttaa atgtttggat ggtntgttgt tctgcctgga 300
gacaaagggt ctaacataga tttaagttga ataacattaa cgggtgctaaa aaatgaaaaa 360

```

ttctaaccga agacatgaca ttcttagctg taa

393

<210> 1630

<211> 317

<212> DNA

<213> Homo sapiens

<400> 1630

```
ctgcaagaat atcagaaatc aatacaaaca agtattgaca ggtgttacag acatgcaaaa 60
tatccttcaa tgcaacgaat ttttaagaaa tcagctagcc tatattaatc agatgtttta 120
ggtcaaacca agtttccatc tcgggctcag tgaaatagta ttaactcatt gagtctcctt 180
tccccagga atgttgggaa tggcagaaca gaaagagcta tcaactcctta aattctttta 240
tgcgagtgtt actccaacac ttattttact tggtttactt ggaatgtatg agaggaaact 300
gatgtttttt acaatgg                                     317
```

<210> 1631

<211> 262

<212> DNA

<213> Homo sapiens

<400> 1631

```
ccttaggcaa gtcaccttac ttatctaaga ctgtttcccc acctggaaga tgccctacaa 60
gcctcctgtg gctgtgttta gaaagcatgc ccggcctttc ttgacagcca gccacccag 120
atgatggcag ggcaaggaag actgtagga gtcagagtgc tccccacagg tggaaggaaa 180
ctgggccaac tctactttgt aagccatagg gtgccaggta gcccggccac cctgagcctg 240
tgcttcact gccccgcgt gg                                     262
```

<210> 1632

<211> 138

<212> DNA

<213> Homo sapiens

<400> 1632

```
ctggaattaa ttcttcgaca actccagacc gaccttcgga agggaaaaca agacaaggcc 60
gttctccaag cagaagtgca gcacctgaga caggacaaca tgagactgca ggaggagtcc 120
cagaccgcga cagctcag                                     138
```

<210> 1633

<211> 192

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(192)

<223> n = A,T,C or G

<400> 1633

```
ccttgaaggg acctcanagc aaaggaagag acctgggtgt ggtgaggcat cccanggcac 60
ggaagggacc ggttgtgctn ngggaatcca ctgnncctc cttggnnaaa aaagcacaac 120
acatcataca tatttaccag accagaagcg ctggcccaa gtctcccaa cctggtcggg 180
ggaacctcct gg                                     192
```

<210> 1634

<211> 447

<212> DNA

<213> Homo sapiens

<400> 1634

```
ctgcttttaa aggtcttaaa tcaactgaat accttgactt gagcttcaat cagatagcca 60
gactgccttc tggctcctct gtctctcttc taactctcta cttagacaac aataagatca 120
gcaacatccc tgatgagtat ttcaagcggt ttaatgcatt gcagtatctg cgtttatctc 180
acaacgaact ggctgatagt ggaatacctg gaaattcttt caatgtgtca tccctgggtg 240
agctggatct gtcctataac aagcttaaaa acataccaac tgtcaatgaa aaccttgaaa 300
actattacct ggaggtcaat caacttgaga agtttgacat aaagagcttc tgcaagatcc 360
tgggggcatt atcctactcc aagatcaagc atttgcgttt ggatggcaat cgcactctcag 420
aaaccagtct tccaccgat atgtatg 447
```

<210> 1635

<211> 364

<212> DNA

<213> Homo sapiens

<400> 1635

```
gtttttatttg agacataaaa acacatgtgt ttctattaca tagtgtgggg tttaggggcc 60
tggtttctaa gacaagactt tatttcaccc tgtatcacag cttcctggga aatgaattag 120
ggagcaagag acggcctggc aagaaaatca ttattgttgc tgggaagttg caaagaaagg 180
ggagagttta ttcaaattag tgtaacagag cccccaggat gaagagagtg gtgcagggaa 240
aaggtctaaa ttcttggtgt tgggtggggac actggcacat cccacagcaa ggactcagcc 300
ctcaacggcg gcggctgggt cttgggaggg gagtgggtggg agggtaaggg ctctcagct 360
ccct 364
```

<210> 1636

<211> 399

<212> DNA

<213> Homo sapiens

<400> 1636

```
ctggctggct agactgtttg tgcgccaaga ggatggtcag cgctgctttc cagcctggct 60
ctgctggggc gctggcatct ggttcagttc caccattctc cctgctttct ttgccaagt 120
tgatattcac ccaagggcac cagtctctat gctgagaggt gggatcaaag aagcttcggg 180
aagatgtgtc cgaactgctg gaggagcaga ggcgagctcg cttggctttc cgcagagggc 240
tagatggtac ctccaggcca ggggtgtctc ctgttcccat gcttcgggtc actgggagag 300
ttctgggtgt ggggctagca gcctctggct caggacgggt aacaggactg gaagagtccc 360
agctccgagt tcgagagaca atgggaccag ggctctttt 399
```

<210> 1637

<211> 246

<212> DNA

<213> Homo sapiens

<400> 1637

```
ctgagctttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa 60
agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatata ttaacaaagc 120
aatagctctc aagcagcaga gcatctcgag gaagaaagct tgcccggtcg ccatcccatc 180
atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa 240
agcttg 246
```


<210> 1638
 <211> 453
 <212> DNA
 <213> Homo sapiens

<400> 1638
 ccaagagttc tccactgtga agactgaaag gacctggtga catttcggca tcagtcctgt 60
 taccacttgg aggtaacaga agcaggctcg tgtcctcctt taattctacc acactacatg 120
 actcgcaatt ggttctgaaa ttagaacgtt caccatcgta cttaaaatct taggggcatg 180
 aagagtcagc tagaacaagg aaaaagaaag tcgcaggtag taggtaagta ggtgggcaca 240
 tgaaaagcca agctgctctg tccaacacca gtgtacatgt gctttaacta aatgaactcc 300
 agaggccaac agcagcagac ctgctcaatt caccttccaa atcagaacaa gacaaaaag 360
 ctcaggcttg agttgtcaac tatgcatagg ttccgccagt gatgaggagc tcgtaagcag 420
 gatctctact ccttctgcac aacacgatgc aag 453

<210> 1639
 <211> 197
 <212> DNA
 <213> Homo sapiens

<400> 1639
 tttgctgttc gtgatatgag acagacagtt gcggtgggtg tcatcaaagc agtggacaag 60
 aaggctgctg gagctggcaa ggtcaccaag tctgccaga aagctcagaa ggctaaatga 120
 atattatccc taatacctgc caccctcctc ttaatcagtg gtggaagaac ggtctcagaa 180
 ctgtttgttt caattgg 197

<210> 1640
 <211> 278
 <212> DNA
 <213> Homo sapiens

<400> 1640
 ccagagcggg gagtcccacc acctcgaact ctgggaattc gagccacagc tctgccagta 60
 cccaagact cagcactagt ctgatgacct gctaattcac tgacagcata gggctgtctg 120
 ttgtttttgc gcaagttggg gtgaacaaag ttcacaatat ctggtcgaat aggagccttg 180
 aatacagcag gcaaagtgc atttttgcca gatgactccc ccttttcgga gtacaccgat 240
 atcagtgggc gagcgcacgc catggcggac ctcggccg 278

<210> 1641
 <211> 227
 <212> DNA
 <213> Homo sapiens

<400> 1641
 ccattgttcc cgtgcatcga agcttgacag cagcttcagg tcctcggtaa acataactct 60
 ctgggggtggc ttgggcccac ccaggaaggt accacatagc ctcttcaagt agctcatgtc 120
 cacgtttaga aagttgtgcc cggcttgcca cgtggtattc cgtttgttga catagttgac 180
 cagctcatcc gacaggggat ggaaagaggg cctgctccgg gcattgg 227

<210> 1642
 <211> 299
 <212> DNA
 <213> Homo sapiens

<223> n = A,T,C or G

<400> 1646

```
ctgtggccgg attgatgggg cccccacttc ctagggctga aggcaagttg aaggaagcag 60
caggagtacc ggaatgaaaa ccttgtttct caaaggactg ctgggttttg gagtacacag 120
aacccgagat atctggcacg cccgtgttac tggagggtgac tgaaacacca gtgttgatc 180
catgagaccc atatccactc ggctgttgga aaggggtggc cgatgcattc aactgacat 240
tcacaccatg ctgcttgga gaggtaggag ccacagggaa cacagcaggc ccatactgga 300
aggtgctggg gagggccggg acccctgtat agtatggcag gctggtgtaa actgtagcca 360
ggaggcagcg ccgggttcag gaatgtctgc tgcgtggnat ggtgagtctg cgtctggttt 420
ctgttggggg ttg 433
```

<210> 1647

<211> 451

<212> DNA

<213> Homo sapiens

<400> 1647

```
ccagcttgca agcacgctgg caaatctctg tcaggtcagc tccagagaag ccattagtca 60
ttttagccag gaactccaag tccacatcct tggcaactgg ggacttgccg aggttagcct 120
tgaggatggc aacacgggac ttctcatcag gaagtgggat gtagatgagc tgatcaagac 180
ggccaggtct gaggatggca ggatcaatga tgtcaggccg gttggtagcg ccaatgatga 240
acacattttt ttttgtggac atgccatcca tttctgtcag gatctggttg atgactcgg 300
cagcagcccc accaccatct ccaatgttac ctccacgagc cttggcaatc gaatccagct 360
catcaaagaa tagcacacag ggggcagctt ggcgggcctt gtcaaagatt tctctgacat 420
tggcctcaga ctccccaac cacatggtga g 451
```

<210> 1648

<211> 176

<212> DNA

<213> Homo sapiens

<400> 1648

```
cctaaacgag gatttcagct tccattatgc ccaactccag tccaacatca ttgaggcgat 60
taatgagctg ctagtggagc tggaaggagc aatggagaac attgcagccc aggctctgga 120
gcacattcac tccaatgagg tgatcatgac cattggcttc tcccgaacag tagagg 176
```

<210> 1649

<211> 435

<212> DNA

<213> Homo sapiens

<400> 1649

```
tgtggctgtg ccgttggtcc tgtgcggtca ctagccaag atgcctgagg aaaccagac 60
ccaagaccaa ccgatggagg aggaggaggt tgagacgttc gcctttcagg cagaaattgc 120
ccagttgatg tcattgatca tcaatacttt ctactcgaac aaagagatct ttctgagaga 180
gctcatttca aattcatcag atgcattgga caaaatccgg tatgaaagct tgacagaccc 240
cagtaaatta gactctggga aagagctgca tattaacctt ataccgaaca aacaagatcg 300
aactctcact attgtggata ctggaattgg aatgaccaag gctgacttga tcaataacct 360
tggtactatc gccaaagtct ggaccaaagc gttcatggaa gctttgcagg ctggtgcaga 420
tatctctatg attgg 435
```

<210> 1650

<211> 246

<212> DNA

<213> Homo sapiens

<400> 1650

```
ccatgtctgt attgtaactg gtaaaaggct tcaagtcaga ttgatgatca agaaaagtca 60
aaaccccagc ccaagattgg gaaagcaggt ggtggttcca agctttttaa aaattattga 120
agctctccat cctgttctgt gagtgtgtct tctctttctc cttcacgtca tagccgtgac 180
ccaccgttca tctctgtctt tgcgtaaaaga tgaccgatgg agtccaaagc caagtggctt 240
caccag                                           246
```

<210> 1651

<211> 400

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(400)

<223> n = A,T,C or G

<400> 1651

```
cggcaagttc tcccaggaga aagccatggt cagttcgagc gccaaagaccg tgaagcccaa 60
tggcgagaag ccggacgagt tcgagtcagg catctcccag gctcttctgg agctggagat 120
gaactcggac ctcaaggctc agctcaggga gctgaatatt acggcagcta nngaaattga 180
agttggtggt ggtcggaaaag ctatcataat ctttgttccc gtccctcaac tgaaatcttt 240
ccagaaaatc caagtccggc tagtacgcga attggagaaa aagttcagtg ggaagcatgt 300
cgnctttatc ggctcagagg aggaattctg cctaagccaa ctcnaaaaag ccgnacnaaa 360
aattanngca aaaagcgtnc caggagccgt nctctgacag                                           400
```

<210> 1652

<211> 338

<212> DNA

<213> Homo sapiens

<400> 1652

```
ctggggggtgc ccatcttctg tgctctgtgg tacatatctg tgtcgccaaa gtagcgtgcc 60
cggtagacga agccttcctt ctgctgcttc tcttccagc agttgttccg gaggttggcg 120
atataatcat ctccacatt ccgctcgact gttttgaggc tggagcctgt gtactcttcg 180
gagaaaagtgt ctccacata gtagacgaca cccagggtgt cagtgaactg cctgtggatg 240
tggcccacag acggtcttgg actcagactg tagggtggac tggagaccat gagctggctg 300
agagctgaca cgagaatcag gatgaggata ggcacacag                                           338
```

<210> 1653

<211> 167

<212> DNA

<213> Homo sapiens

<400> 1653

```
gcggtggagc cgccaccaa atgcagattt tcgtggaaac ccttacgggg aagaccatca 60
ccctcgaggt tgaaccctcg gatacgatag aaaatgtaaa ggccaagatc caggataagg 120
aaggaattcc tcttgatcgg cagagactga tctttgctgg caagcag                                           167
```

<210> 1654
 <211> 1034
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(1034)
 <223> n = A,T,C or G

<400> 1654
 atgcatgctc gagcgggccgc cagtgtgatg gatatactgca gaattcgccc ttagcgtggg 60
 cgcgggccgag gtccaagagg gagataaanac aaactttctca aacaaaaaga aaagaaaaac 120
 gaatgattca tctgctttta tcagtgtgat taatgcagca cccattgccc cggaaccgt 180
 ttctgctgta ctatctggat actaaaatgt tacggaagta gctctttgtt ctccctcact 240
 ctgcccttag ttaatagaaa ttcagactcg ccaagtaagg ctttgtgcat agtgtcttca 300
 tgtcgcgtat agttgagcgc gttcttagca gttggcttca tggacagctc attagtgttt 360
 tgacttttct taccagcgt taattgaatt cttgctttta gacaacttcc tttttgtagt 420
 ggtgaacctt gcccttttagt acagttcaag tgaatctgga taattgttca tctttgcttt 480
 agcttagata ccatgtagtg gtctgtggct acaggaagct ggttctgtct gcttcacag 540
 tctgcttaaa aaactgtctg acttcgtgaa tatagagacc aagtttacca cttctgatga 600
 agagaccaat taagattcat tcctcattct gtttctttcc agtgggagaa gagtcccat 660
 gaaataagat gaaactgatt ccatgcacta gtacatgtag gcttctccct tgcgcaaagc 720
 ttaacaattt gtaggaaact ttgggtcttt ttgtcccaag aaaaaggaat gtcttgacag 780
 gcttaaagct ttctgctccc ttgcacctta aaactcgaaa gttagnaaa atccctttaa 840
 agggcttttt ttaatagcca gaacttccca aaaggaatgg cnttttaggg aatttcntag 900
 ccatngcttt ttaaatttaa agaaattttt aanaaccttg cccnnggggn ggggnccgc 960
 tccaaaaagg gnggnaaaaa ttcccagcc nacctttng gggggggccn cgttttctt 1020
 tnnngggggg aanc 1034

<210> 1655
 <211> 487
 <212> DNA
 <213> Homo sapiens

<400> 1655
 atgcatgctc gagcgggccgc cagtgtgatg gatatactgca gaattcgccc ttctgagcgg 60
 ccgcccgggc aggtcctact cttctccgtc cattgtacta tctgcccgtg gtggggatgg 120
 cagtaggac atatttgatg acttccgaga agcatattat tggctccgtc ataatactcc 180
 agaggatgcg aaggtcatgt cctggtggga ttatggctat cagattacag ctatggcaaa 240
 ccgaacaatt ttagtgagca ataacacatg gaataatacc catatttctc gagtagggca 300
 ggcaatggcg tccacagagg aaaaagccta tgagatcatg agggagctcg atgtcagcta 360
 tgtgctggtc atttttggag gacctcggcc gcgaccacgc taaggcgaa ttccagcaca 420
 ctggcgcccg ttactagtgg atccgagctc ggtaccaagc ttggcgtaat catggtcata 480
 gctgttt 487

<210> 1656
 <211> 514
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(514)

<223> n = A,T,C or G

<400> 1656

```
atgcatgctc gagcgggccc ccagtgtgat ggatatctgc agaattcgcc cttancgtgg 60
tcgcgggcca ggtcctaccc ataatccaga gaggcttgcc cagaggagga ctacgtgggg 120
gacgtgccac cagaacccta cttgggggcg ggatgtcact ccgagggtcaa aacctgctcc 180
gaggtggacg agccgtagct ccccgaaatg gcttaagaag aggtggtggt cgaggctcgtg 240
gaggtcctgg gagagggggc ctagggcgtg gagctatggg tcgtggcgga atcgggtggt 300
gaggtcgggg tatgataggt cggggaagag ggggctttgg aggccgaggc cgaggccgtg 360
gacgagggag aggtgccctt gctcgccctg tattgaccaa ggagcagacc tgcccggggc 420
gccgctcgaa gggcgaattc cagcacactg gcggccgtta ctagtggatc cgagctcggt 480
accaagcttg gcgtaatcat ggtcatagct gttt 514
```

<210> 1657

<211> 605

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(605)

<223> n = A,T,C or G

<400> 1657

```
atgcatgctc gagcgggccc cagtgtgatg gatattctgca gaattcgccc tttcgagcgg 60
ccgcccgggc aggtccanac gctgacattg nttctgagtc cttaagcagg aaggatttga 120
aatcctggag cttggcagtc ttgctcttca cctctaagcc aatgttgacc ccttcacta 180
taaagtcac aactctccgg aagtcacct caccggaactg tcgagaagtt aaggctgggg 240
ccccaagccg caggccgccc ggtgtgatgg cacttcggtc tccaggacag gtgttcttgt 300
tggcagtgat ggatacaagc tctagcacc gctcagccc agctccatcc aggcccttgg 360
gccgcaggtc caccagcacc aggtggttgt cagtaccacc tgataccagt gagtgcctc 420
gccctagcag ggcattctgc atggcccag cattcttcag aacctgcagg gagtactccc 480
ggaacatggg ggtgcaggac ctcgcccgcg accacgctaa gggcgaattc cagcacactg 540
gcggccgtta ctagtggatc cgagctcggt accaagcttg gcgtaatcat ggtcatagct 600
gtttc 605
```

<210> 1658

<211> 784

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(784)

<223> n = A,T,C or G

<400> 1658

```
agnnttcgn cgccctcna gntgcatgct cgagcgggccg cgcagtgaga tgnatatctg 60
cagaattcgc cttancgtg ggcnangca tgacgctcgg gatcagaact aaaacaagt 120
agatcacccc tctaattatt tctgaactng gttataaaaa gcttataaga tttttatgaa 180
gcancactg tatgatattt taagcaaata tggtatttaa aatattgatc cttcccttgg 240
accaccttca tgtagttgg gtattataaa taagagatac aacctgaat atattatgtt 300
tatacaaaat caatctgaac acaattcata aagatttctc ttttatacct tcctcactgg 360
ccccctccac ctgcccatag tcaccaaatt ctgttttaaa tcaatgacct aagatcaaca 420
```

```

atgaagtatt ttataaatgt atttatgctg ctagactgtg ggtcaaagt ttccattttc 480
aaattattta gaattcttat gagtttaaaa tttgtaaatt tctaaatcca atcatgtaaa 540
atgaaactgt tgctccattg gagtagtctc ccacctaaat atcaagatgg ctatatgcta 600
aaaagagaaa atatgggtcaa gtctaaaatg gctaattgtc ctatgatgct attatcatag 660
actaaccgac atttatcttc aaaacaccaa attgtcttta gaaaaaatta atngtgatta 720
ccaggtagaa ggacctgccc gggcggnccg ctcgaaaggg ccgaaattcc agccccacct 780
gggc

```

<210> 1659

<211> 789

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(789)

<223> n = A,T,C or G

<400> 1659

```

tngngccctc tagatgcang ctcgagcggc cgccagtgtg atggatatct gcagaattcg 60
cccttagcgt ggtecgggcc gaggtccatt aaagataagt ttggctaact attttactga 120
agagactaat ggtcttcctt ctgttgtaact gctatgtttc ttgatctgtt tttccccaat 180
gtaacagtct acattgaagt cctttagctc tctccatata ctaattgaca tttgttaagg 240
attcaatatt ttgtgaattc tttttaccct taaaatgcat atctttcaga gagataagaa 300
tgaattttgc aataatttat atgcagagtg tgcttatggg tttctgggag ttcaagttag 360
taccgccagag tgcttaaaag tacgatgcta aattctaagg ctaatgtaat gactgtagat 420
tatctatgtc cacattgttc aacagaaata taatgtgaac cacaacataa tttttaattt 480
tctagtagcc atattaaaaa agaaacaagc aaaattaatt ttaataacag tttatgtaac 540
ccagtatatt aaaaatatca tttcaacatg taatcaatat aaaagattat taatgaaaca 600
ccttatcctc tttttcttcc atgctaagtc ttagatttga gtgtattttg cactcacagc 660
acatctcaat tctgactgga cctgccggg cggcgcgtcg aaagggcgaa ttccagcaca 720
ctgggcggcc gttactagtg gatccgagct ccggtacca gcttggcgta atcatggtca 780
tagctgttt

```

<210> 1660

<211> 559

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(559)

<223> n = A,T,C or G

<400> 1660

```

ccnccgctc tagatgcatg ctcgagcggc cgccagtgtg atggatatct gcngaattcg 60
ccctttccag cgcccgcccg ggcaggtcca tcagacttct tgggtgcctg gctatattca 120
atgtgaagta aaaaatatcc caagtcttac accaaaatag aggctctgac ttagaagtat 180
gcttttagct ttctttttaa ataagacatt ctggaagaaa aaaaaagaaa aaggaaagaa 240
aatcaagttt gaaacacagt taacacttat tttggcaaga aagcaaccaa aatctaaaaa 300
gcataaacta tngtccaaa tgnaaaaggn attacagaac aaactgcaag aggggaaaat 360
taaagccnca ctgaacgaaa aaatacagta tgtctaacat tttggaattg naattttaa 420
cctaagggca aaagctgaaa aatcatgctt anacctnggn cnggaccacn ctaagggcga 480
attccancac actggcggn cgttactagt gatccnanc cgggtaccaag cttggcgtaa 540

```

tcctnnggcat agctgtttc

559

<210> 1661

<211> 453

<212> DNA

<213> Homo sapiens

<400> 1661

```

ttgggccctc tagatgcatg ctcgagcggc cgccagtgtg atggatatct gcagaattcg 60
cccttttcgag cggccgcccc ggccaggtctg cagtgtccct tttatatca tgctagtgtt 120
gagacatact tgactaactt gggaacagtt cgatatattg acaaccgtca acttaagaaa 180
atcaacagct tttggcccca gcgtccaagt gaacttttca tggagtgcag aatctcaaat 240
ggacaaaata ctttgtcttt ttaaatactg aaaatttaaat tattagtact atgactgaaa 300
gattcttcat ggctaaaaag ctctgcatca aactcaattc aggaggacct cggccgcgac 360
cacgctaagg gcgaattcca gcacactggc ggccgttact agtggatccg agctcggtag 420
caagcttggc gtaatcatgg tcatagctgt ttc 453

```

<210> 1662

<211> 809

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(809)

<223> n = A,T,C or G

<400> 1662

```

ctcgagcggc cgccantgtg atggntatct gcagaattcg cccttanccg ccgcccgggc 60
aggtccttag ccaaagaatg cagtggagcc ttccccnngg ggctgcattg tgaatgaata 120
ccaattgaca gcataaaaat taatagtccc atatcagatc tgggaaggggt ttctggggct 180
gtctgatgtc cctatcctgt tgtagtgaac acaatagcag aaaattcttt ctgggtccat 240
ctgctataaa gtcttggtta aacagcatta ctatgaagag gatgaactca cctaccttca 300
natggaggaa aagtgaagaa gacttaggct ttagtctctc atgacttttc ttaagcacta 360
cctacctgta ataagctgag tgcaaaagga tgccgaagaa aatctgcacc cagaagctgt 420
tagaaagcac tgcaangaa cagggnatga ataaaataaa nagntcttaa taaaccctta 480
agattctttg ntcaaggggn actttgccaa aaggggcaga atangngggg aaagagttgc 540
ttttaatcta gctctacact ggcntttgaa aataaaaattt gcccattnng aaatatatng 600
ggntataatt aaaatgnngc tttttacact gnggggggcta tataaaaact gggtagnaaa 660
atttccaccg agcatntatg gngatttgnt cacagnaaac ctccgggcng gaccacgct 720
aagggnggaa ttccagcnac antggggggg ncngntacct anagtggatc ccnagnctng 780
gggnccccna anctttgggg gngtnaatc 809

```

<210> 1663

<211> 585

<212> DNA

<213> Homo sapiens

<400> 1663

```

ttgggccctc tagatgcatg ctcgagcggc cgccagtgtg atggatatct gcagaattcg 60
cccttgccgc ccgggcaggt gatggatgag gagcaaaaac tttatacgga tgatgaagat 120
gatattctaca aggctaataa cattgcctat gaagatgtgg tggggggaga agactggaac 180
ccagtagagg agaaaataga gagtcaaacc caggaagagg tgagagacag caaagagaat 240
atagaaaaaa atgaacaaat caacgatgag atgaaacgct cagggcagct tggcatccag 300

```



```

gaagaagatc ttcggaaga gagtaaagac caactctcag atgatgtctc caaagtaatt 360
gcctatttga aaaggttagt aaatgctgca ggaagtggga gggtacagaa tgggcaaaat 420
ggggaaaggg ccaccaggct ttttgagaaa cctcttgatt ctcagtctat ttatcagacc 480
tcggccgcga ccacgctaag ggcgattcc agcacactgg cggccgttac tagtggatcc 540
gagctcggta ccaagcttgg cgtaatcatg gtcatactg tttcc 585

```

<210> 1664

<211> 999

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(999)

<223> n = A,T,C or G

<400> 1664

```

anccngetcn agcggccgcc antgtgatgg atatctgcag aattcgccct ttcgagcggg 60
ccgcccgggc aggtctgaca atngattaaa caggcgacat gcaaccccca ctaagggttaa 120
aagtccaaaa ctactcacac gcatctcttn attggggaaa agctgagact attatncatt 180
cttggtagnc ttgcaacctt gcatgaagag caccatttgc atttctttca tctttcagaa 240
agcaccggta tctgttccaa ggnnctaaca gtacnaaaat acnttntggg attacacctt 300
tnaaacccaa nactgttntc attaaaaata attttggnnt gtaacaaaat tatgaaatac 360
aatgcaagca cctnggtata gcattattac tgaaaccact taattcccag ctttttgagt 420
tttttaaaaa aaccactgc actaagattc acaattcatt gctacatata aattaaagct 480
agtaagaaca cactaacgtc acaagtttct cattctaaag tgcnaaancc ntaatngtct 540
ngaaagtgga acaggggtaa agggcaaaaa ttaacccccc ccacccaat taaagtttcc 600
tggaangtca ntantntttt naatcccaa aggnnncatt tctnttttaa aaaattggnt 660
acctttggaa ctggggtaaa gnaaaatnag gaacccctgg gnggtttttt ttatnttttc 720
ttnaanccaa cccccaatt ccaccttaa aacccccacc cggggggang ccaaaaangnc 780
cacccttgng gaaacncttt tngtgggggn cccggtcgna aaacccaacc nccctntaaa 840
aagggggggg cgnaaaaaaa tttctcccna aganaaaacc acctttgggg cgnggggacn 900
cgntttaccc nttaaaatgg gggaattcc ccgaaagcgt ttggggggtaa ccccaaaaga 960
cctttggggg gggaaaaatg aatgggggnc cattaacn 999

```

<210> 1665

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 1665

gctaaagggtg accccaagaa accaaag

27

<210> 1666

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 1666

ctattaactc gagggagaca gataaacagt ttccttta

37

<210> 1667

<211> 207

<212> PRT

<213> Homo sapiens

<400> 1667

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gln | His | His | His | His | His | His | Ala | Lys | Gly | Asp | Pro | Lys | Lys | Pro |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Lys | Gly | Lys | Met | Ser | Ala | Tyr | Ala | Phe | Phe | Val | Gln | Thr | Cys | Arg | Glu |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Glu | His | Lys | Lys | Lys | Asn | Pro | Glu | Val | Pro | Val | Asn | Phe | Ala | Glu | Phe |
| | | | 35 | | | | 40 | | | | | 45 | | | |
| Ser | Lys | Lys | Cys | Ser | Glu | Arg | Trp | Lys | Thr | Met | Ser | Gly | Lys | Glu | Lys |
| | | | 50 | | | 55 | | | | | 60 | | | | |
| Ser | Lys | Phe | Asp | Glu | Met | Ala | Lys | Ala | Asp | Lys | Val | Arg | Tyr | Asp | Arg |
| 65 | | | | | 70 | | | | 75 | | | | | 80 | |
| Glu | Met | Lys | Asp | Tyr | Gly | Pro | Ala | Lys | Gly | Gly | Lys | Lys | Lys | Lys | Asp |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| Pro | Asn | Ala | Pro | Lys | Arg | Pro | Pro | Ser | Gly | Phe | Phe | Leu | Phe | Cys | Ser |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Glu | Phe | Arg | Pro | Lys | Ile | Lys | Ser | Thr | Asn | Pro | Gly | Ile | Ser | Ile | Gly |
| | | | 115 | | | 120 | | | | | | 125 | | | |
| Asp | Val | Ala | Lys | Lys | Leu | Gly | Glu | Met | Trp | Asn | Asn | Leu | Asn | Asp | Ser |
| | | | 130 | | | 135 | | | | | 140 | | | | |
| Glu | Lys | Gln | Pro | Tyr | Ile | Thr | Lys | Ala | Ala | Lys | Leu | Lys | Glu | Lys | Tyr |
| 145 | | | | | 150 | | | | | 155 | | | | 160 | |
| Glu | Lys | Asp | Val | Ala | Asp | Tyr | Lys | Ser | Lys | Gly | Lys | Phe | Asp | Gly | Ala |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Lys | Gly | Pro | Ala | Lys | Val | Ala | Arg | Lys | Lys | Val | Glu | Glu | Glu | Asp | Glu |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Asp | Glu | |
| | | | 195 | | | | 200 | | | | | | 205 | | |

<210> 1668

<211> 636

<212> DNA

<213> Homo sapiens

<400> 1668

| | | | | | | |
|------------|------------|-------------|-------------|------------|-------------|-----|
| catatgcagc | atcaccacca | tcaccacgct | aaaggtgacc | ccaagaaacc | aaagggcaag | 60 |
| atgtccgctt | atgccttctt | tgtgcagaca | tgacagagaag | aacataagaa | gaaaaaccca | 120 |
| gaggtccctg | tcaattttgc | ggaattttcc | aagaagtgtc | ctgagaggtg | gaagacgatg | 180 |
| tccgggaaag | agaaatctaa | atttgatgaa | atggcaaaag | cagataaagt | gcgctatgat | 240 |
| cgggaaatga | aggattatgg | accagctaag | ggaggcaaga | agaagaagga | tcctaattgct | 300 |
| cccaaaaggc | caccgtctgg | attcttctctg | ttctgttcag | aattccgccc | caagatcaaa | 360 |
| tccacaaacc | ccggcatctc | tattggagac | gtggcaaaaa | agctgggtga | gatgtggaat | 420 |
| aatttaaagt | acagtgaaaa | gcagccttac | atcactaagg | cggcaaagct | gaaggagaag | 480 |
| tatgagaagg | atgttgctga | ctataagtgc | aaaggaaagt | ttgatggtgc | aaagggtcca | 540 |

gctaaagttg cccggaaaaa ggtggaagag gaagatgaag aagaggagga ggaagaagag 600
gaggaggagg aggaggagga tgaataatga ctcgag 636

<210> 1669

<211> 2821

<212> DNA

<213> Homo sapiens

<400> 1669

```
ccacgcgtcc ggcgcgcgcg gcgcagggga ggcgagaggc gccccccggt ggagagcctg 60
agccccgcgc aagtctggcg gcacctggcg agcggagccg gagtcgggct ggggaccgcg 120
gggttgaggc cggaccgcgg cggggtcggg ggagaaacgc gcgctgccct ggcacgggcc 180
ccaaccccc ggccgcgcgg aatggtatgg cccggccgga gttaaggccg gggggaggcg 240
gcgagtcctc cggcggcggc gacgatgggg ctgcgtgcag gaggaacgct gggcagggcc 300
ggcgcggggtc gggggggcgcc cgagggggcc gggccgagcg gcggcgcgca gggcggcagc 360
atccactcgg gccgcctcgc cgcggtgcac aacgtgccgc tgagcgtgct catccggccg 420
ctgccgtccg tgttggaacc cgccaagggt cagagcctcg tggacacgat ccgggaggac 480
ccagacagcg tgccccccat cgatgtcctc tggatcaaag gggcccaggg aggtgactac 540
ttctactcct ttgggggctg ccaccgctac gcggcctacc agcaactgca gcgagagacc 600
atccccgccca agcttggtcca gtccactctc tcagacctaa ggggtgtacct gggagcatcc 660
acaccagact tgcagtagca gcctccttgg cactgtctgc caccttcaag agcccagaag 720
acacacctgg cctccagcag gctgggccat gcagaaggga tagcaggggt gcattctctt 780
tgcacctggc gagaggggtc gactctgggc accctctca ccggtacaa ggccttgga 840
tcaactgtaca gtgtgggagc ccagttccc acctctgtga caataggatc atggccttac 900
ccttgaagca ttaccgagaa ggagaacaga gatgggcttg aagagccacg tgctgcccgc 960
tccaaattcc caaggacaag gatccctctg cattttgtc tatgtaacct cttatatgga 1020
ctacattcag ctgcaaggaa aggaaaacct tgattgcagt ggtttaaca aacagaagat 1080
tgtttttcca catagcatgg attctggaga tgggtggcta atggtattgg ttcaacaact 1140
ccacgaagggt aggggtcacg tcttggtacc ttttgctta atctcagtgc tcgttacttc 1200
atggtcccaa gatggctgct gtatcccaa gaatcatgtc tgcgttcaag gaaggagggg 1260
tgaggaaga ggaaggcca aactagctgg acccgtcacc ttctatcaga aagtaaaacc 1320
tcgtcagaag tctgtttcct gctctctccc tctgcatac ttcaactaga tgcccttgga 1380
ccgagccagc taccattgca cctctagctg caaacaagc taagacagca gggaacagaa 1440
ttgtcatggc tgaatagacc aatcgtgttc catctactga gactggcaca ctgcctcctg 1500
caataaaaact gggatcccat taccaagaga gaaatgcaga attgtgtacc agtttagctt 1560
tgctgtgtaa caaacatcc ccaaacttgg cagctagaaa caaacctgt attttccac 1620
aatcctatgg gttggcaatt tgggctgggc tcaacagggc agttctgctg ctcacacctg 1680
ggatccctca tggagctaag gtcagctgtt acctcagctg ggctggatg gtctaggata 1740
gccttactca cttgcctggc aggtgacagg ctgttggtg gaattgctt gttctcctcc 1800
atgtggcctc tccagcaggc tagctcaggc ttattcacat gatggcttca ggattccaaa 1860
gagagtgaga gtagaagctg aaagacttct tgagttcttg gcctggaact gggactagga 1920
cagtgtcact tctgctaagt tcttttggtc agagcaaata ccaaggctt acccagattc 1980
aagggatgag aaacagacta catgtcttga tgaggggaac cacaagagc ttgtggccat 2040
ttttcaccta tcacaaataa ttttggtagg gtattttatt ggataaagg atttccctct 2100
tcccccttct tctctgtctc atggggcctc actctgccaa gttggaaggc actaagacat 2160
tgtcctggcc ctcagggctc aggggaagag gtgttggggc aggaagttag tctctccatg 2220
ggctggaccc actgtagtag gtagtcctcc ttgtctgcac tgctggtagt ggggttaggc 2280
aggtaggaca ttccagagg gcttctgaaa accaagagtc cctggggaaa ggggaacagag 2340
taaggcaggc cttgtttctc ctgccctcta agggaaactg gtcactcggc acttttaagc 2400
ctcagtttct ccagttcaat aataaggaca agagcttttc ccatgcattc tctttcccg 2460
ggaaagttga ctgaggtgac cagtaataga attgaaaagg gagagtgtct tcagtgaat 2520
gtggcatcct ggattgggtc ttggaacaaa aacaggacat tagtgggaaa attggaatc 2580
tgaaaaaagt ctgaatttta gttaatatat caatttcagt ctcttggttt tgacagatgt 2640
```

```

accatgggtga tgtaagatgt tgaccttggg gtaggctggg tgaagggtat acaggaactc 2700
tttgtactat ctctgcaact tctctgtaaa tctagtatca ttccaaaata aaagttttatt 2760
taatttaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 2820
a                                                                 2821

```

```

<210> 1670
<211> 137
<212> PRT
<213> Homo sapiens

```

```

<400> 1670
Met Gly Leu Arg Ala Gly Gly Thr Leu Gly Arg Ala Gly Ala Gly Arg
      5                      10                      15

Gly Ala Pro Glu Gly Pro Gly Pro Ser Gly Gly Ala Gln Gly Gly Ser
      20                      25                      30

Ile His Ser Gly Arg Ile Ala Ala Val His Asn Val Pro Leu Ser Val
      35                      40                      45

Leu Ile Arg Pro Leu Pro Ser Val Leu Asp Pro Ala Lys Val Gln Ser
      50                      55                      60

Leu Val Asp Thr Ile Arg Glu Asp Pro Asp Ser Val Pro Pro Ile Asp
      65                      70                      75                      80

Val Leu Trp Ile Lys Gly Ala Gln Gly Gly Asp Tyr Phe Tyr Ser Phe
      85                      90                      95

Gly Gly Cys His Arg Tyr Ala Ala Tyr Gln Gln Leu Gln Arg Glu Thr
      100                     105                     110

Ile Pro Ala Lys Leu Val Gln Ser Thr Leu Ser Asp Leu Arg Val Tyr
      115                     120                     125

Leu Gly Ala Ser Thr Pro Asp Leu Gln
      130                     135

```

```

<210> 1671
<211> 109
<212> PRT
<213> Homo sapiens

```

```

<400> 1671
Met Ala Arg Pro Glu Leu Arg Pro Gly Gly Gly Gly Glu Ser Arg Gly
      5                      10                      15

Gly Gly Asp Asp Gly Ala Ala Cys Arg Arg Asn Ala Gly Gln Gly Arg
      20                      25                      30

Arg Gly Ser Gly Gly Ala Arg Gly Ala Arg Ala Glu Arg Arg Arg Ala
      35                      40                      45

```


<400> 1673

Met Asp Tyr Ile Gln Leu Gln Gly Lys Glu Asn Leu Asp Cys Ser Gly
 5 10 15

Leu Asn Lys Gln Lys Ile Val Phe Pro His Ser Met Asp Ser Gly Asp
 20 25 30

Gly Trp Leu Met Val Leu Val Gln Gln Leu His Glu Gly Arg Gly His
 35 40 45

Val Leu Asp Pro Phe Ala Leu Ile Ser Val Leu Val Thr Ser Trp Ser
 50 55 60

Gln Asp Gly Cys Cys Ile Pro Lys Asn His Val Cys Val Gln Gly Arg
 65 70 75 80

Arg Gly Gly Gly Arg Gly Arg Ala Lys Leu Ala Gly Pro Val Thr Phe
 85 90 95

Tyr Gln Lys Val Lys Pro Arg Gln Lys Ser Val Ser Cys Ser Leu Pro
 100 105 110

Leu His Ile Phe Thr
 115

<210> 1674

<211> 90

<212> PRT

<213> Homo sapiens

<400> 1674

Met Asp Ser Gly Asp Gly Trp Leu Met Val Leu Val Gln Gln Leu His
 5 10 15

Glu Gly Arg Gly His Val Leu Asp Pro Phe Ala Leu Ile Ser Val Leu
 20 25 30

Val Thr Ser Trp Ser Gln Asp Gly Cys Cys Ile Pro Lys Asn His Val
 35 40 45

Cys Val Gln Gly Arg Arg Gly Gly Gly Arg Gly Arg Ala Lys Leu Ala
 50 55 60

Gly Pro Val Thr Phe Tyr Gln Lys Val Lys Pro Arg Gln Lys Ser Val
 65 70 75 80

Ser Cys Ser Leu Pro Leu His Ile Phe Thr
 85 90

<210> 1675

<211> 102

<213> Homo sapiens

Met Gln Asn Cys Val Pro Val Ser Phe Cys Cys Val Thr Asn His Pro
5 10 15

Gln Thr Trp Gln Leu Glu Thr Asn Pro Val Phe Ser His Asn Pro Met
20 25 30

Gly Trp Gln Phe Gly Leu Gly Ser Thr Gly Gln Phe Cys Cys Ser His
35 40 45

Leu Gly Ser Leu Met Glu Leu Arg Ser Ala Val Thr Ser Ala Gly Pro
50 55 60

Gly Trp Ser Arg Ile Ala Leu Leu Thr Cys Leu Ala Gly Asp Arg Leu
65 70 75 80

Leu Ala Gly Ile Ala Trp Phe Ser Ser Met Trp Pro Leu Gln Gln Ala
85 90 95

Ser Ser Gly Leu Phe Thr
100

<211> 1336

<213> Homo sapiens

| | | | | | | |
|-------------|------------|------------|------------|------------|------------|------|
| ctctaagcag | catgtaacct | ggcctgcac | caggaaatag | aggacttcgg | atccttctaa | 60 |
| ccctaccacc | caactggccc | cagtacattc | attctctcag | gaaaaaaaaa | aaggtcccca | 120 |
| cagcaaagaa | aaggaatagg | atcaagagat | acgtggctgc | tggcagagca | agcatgaatt | 180 |
| cgatgacttc | agcagttccg | gtggccaatt | ctgtgttggt | ggtggcacc | cacaatgggt | 240 |
| atcctgtgac | cccaggaatt | atgtctcacg | tgcacctgta | tccaaacagc | cagccgcaag | 300 |
| tccacctagt | tcttggaac | ccacctagtt | tgggtgctga | tgtgaatggg | cagcctgtgc | 360 |
| agaaagctct | gaaagaaggc | aaaaccttgg | gggccatcca | gatcatcatt | ggcctggctc | 420 |
| acatcggcct | cggctccatc | atggcgacgg | tctctgtatg | ggaatacctg | tctatttcat | 480 |
| tctacggagg | ctttcccttc | tggggaggct | tgtggtttat | catttcagga | tctctctccg | 540 |
| tggcagcaga | aaatcagcca | tattcttatt | gcctgctgtc | tggcagtttg | ggcttgaaca | 600 |
| tcgctcagtc | aatctgctct | gcagttggag | tcatactctt | catcacagat | ctaagtattc | 660 |
| cccacccata | tgcctacccc | gactattatc | cttacgcctg | gggtgtgaac | cctggaatgg | 720 |
| cgattttctg | cgtgctgctg | gtcttctgcc | tcttgagatt | tggcatcgca | tgcgcatctt | 780 |
| cccacttttg | ctgccagttg | gtctgctgtc | aatcaagcaa | tgtgagtgtc | atctatccaa | 840 |
| acatctatgc | agcaaacc | gtgatcacc | cagaaccggg | gacctacca | ccaagttatt | 900 |
| ccagtgcagt | ccaagcaaat | aagtaaggct | acagattctg | gaagcatctt | tactgggac | 960 |
| ctaagaaga | cctctcctct | ttctgggctt | ccataacc | ggtcgttcct | gttctgacag | 1020 |
| caggagaaac | gtctctccca | ctgttgttac | tctcaccttc | attcttcaat | tcagtctagg | 1080 |
| aaacctatgct | gtttctctat | caagaagaag | acagagattt | taacagatct | ttaaccaaga | 1140 |
| gggactccct | agggcacatg | catcagaca | tatgtgggca | tccagcctct | ggggccttgg | 1200 |
| cacacacaca | ttcgtgtgct | ctgctgcac | tgcctgtgtg | ggttagagga | acaaatatct | 1260 |

```
<210> 1677
<211> 250
<212> PRT
<213> Homo sapiens
```

Met Asn Ser Met Thr Ser Ala Val Pro Val Ala Asn Ser Val Leu Val
5 10 15

Val Ala Pro His Asn Gly Tyr Pro Val Thr Pro Gly Ile Met Ser His
20 25 30

Val Pro Leu Tyr Pro Asn Ser Gln Pro Gln Val His Leu Val Pro Gly
35 40 45

Asn Pro Pro Ser Leu Val Ser Asn Val Asn Gly Gln Pro Val Gln Lys
50 55 60

Ala Leu Lys Glu Gly Lys Thr Leu Gly Ala Ile Gln Ile Ile Ile Gly
65 70 75 80

Leu Ala His Ile Gly Leu Gly Ser Ile Met Ala Thr Val Leu Val Gly
85 90 95

Glu Tyr Leu Ser Ile Ser Phe Tyr Gly Gly Phe Pro Phe Trp Gly Gly
100 105 110

Leu Trp Phe Ile Ile Ser Gly Ser Leu Ser Val Ala Ala Glu Asn Gln
115 120 125

Pro Tyr Ser Tyr Cys Leu Leu Ser Gly Ser Leu Gly Leu Asn Ile Val
130 135 140

Ser Ala Ile Cys Ser Ala Val Gly Val Ile Leu Phe Ile Thr Asp Leu
145 150 155 160

Ser Ile Pro His Pro Tyr Ala Tyr Pro Asp Tyr Tyr Pro Tyr Ala Trp
165 170 175

Gly Val Asn Pro Gly Met Ala Ile Ser Gly Val Leu Leu Val Phe Cys
180 185 190

Leu Leu Glu Phe Gly Ile Ala Cys Ala Ser Ser His Phe Gly Cys Gln
195 200 205

Leu Val Cys Cys Gln Ser Ser Asn Val Ser Val Ile Tyr Pro Asn Ile
210 215 220

Tyr Ala Ala Asn Pro Val Ile Thr Pro Glu Pro Val Thr Ser Pro Pro
225 230 235 240

Ser Tyr Ser Ser Glu Ile Gln Ala Asn Lys
 245 250

<210> 1678
 <211> 177
 <212> PRT
 <213> Homo sapiens

<400> 1678
 Thr Arg Pro Arg Arg Ala Ala Gln Gly Arg Arg Glu Ala Pro Pro Gly
 5 10 15
 Gly Glu Pro Glu Pro Arg Ala Ser Leu Ala Ala Pro Gly Glu Arg Ser
 20 25 30
 Arg Ser Arg Ala Gly Asp Arg Gly Val Glu Ala Gly Pro Arg Arg Gly
 35 40 45
 Arg Gly Arg Asn Ala Arg Cys Pro Gly Thr Gly Pro Asn Pro Pro Ala
 50 55 60
 Ala Arg Asn Gly Met Ala Arg Pro Glu Leu Arg Pro Gly Gly Gly Gly
 65 70 75 80
 Glu Ser Arg Gly Gly Gly Asp Asp Gly Ala Ala Cys Arg Arg Asn Ala
 85 90 95
 Gly Gln Gly Arg Arg Gly Ser Gly Gly Ala Arg Gly Ala Arg Ala Glu
 100 105 110
 Arg Arg Arg Ala Gly Arg Gln His Pro Leu Gly Pro His Arg Arg Gly
 115 120 125
 Ala Gln Arg Ala Ala Glu Arg Ala His Pro Ala Ala Ala Val Arg Val
 130 135 140
 Gly Pro Arg Gln Gly Ala Glu Pro Arg Gly His Asp Pro Gly Gly Pro
 145 150 155 160
 Arg Gln Arg Ala Pro His Arg Cys Pro Leu Asp Gln Arg Gly Pro Gly
 165 170 175
 Arg

<210> 1679
 <211> 42
 <212> PRT
 <213> Homo sapiens

<400> 1679
 Leu Val Cys Cys Gln Ser Ser Asn Val Ser Val Ile Tyr Pro Asn Ile

